

Supporting Information for “Tautomerization in the Formation and Collision-Induced Dissociation of Alkali Metal Ion-Cytosine Complexes” by Zhibo Yang and M. T. Rodgers*

Table 1S. Vibrational Frequencies and Average Vibrational Energies of Cytosine and M⁺(cytosine)^a

Species	E _{vib} , eV	Vibrational Frequencies ^a , cm ⁻¹
C ₁	0.14 (0.01)	-508, -51, 125, 246, 301, 364, 470, 493, 527, 553, 616, 646, 709, 765, 822, 893, 957, 1049, 1097, 1180, 1283, 1340, 1413, 1510, 1587, 1640, 1751, 1847, 3280, 3313, 3658, 3689, 3800
C ₂	0.14 (0.01)	-376, 179, 212, 329, 430, 481, 483, 543, 545, 581, 650, 725, 750, 771, 916, 965, 979, 1071, 1111, 1237, 1296, 1331, 1394, 1456, 1509, 1595, 1610, 1651, 3110, 3146, 3519, 3585, 3654
C ₃	0.14 (0.01)	-389, 178, 215, 328, 432, 467, 483, 529, 539, 588, 646, 724, 750, 771, 924, 963, 982, 1068, 1109, 1229, 1301, 1342, 1393, 1445, 1506, 1603, 1608, 1649, 3107, 3146, 3522, 3590, 3655
C ₄	0.15 (0.01)	106, 136, 341, 361, 494, 496, 519, 550, 610, 642, 686, 712, 752, 815, 868, 953, 965, 1078, 1135, 1204, 1299, 1375, 1386, 1421, 1484, 1623, 1701, 1797, 3145, 3166, 3388, 3489, 3535
C ₅	0.15 (0.01)	110, 134, 346, 359, 490, 496, 517, 542, 567, 623, 692, 734, 758, 823, 871, 959, 961, 1061, 1134, 1193, 1280, 1376, 1392, 1413, 1475, 1634, 1703, 1794, 3153, 3187, 3343, 3486, 3536
C ₆	0.14 (0.02)	-424, 129, 186, 326, 330, 425, 497, 524, 577, 582, 637, 692, 734, 775, 907, 948, 1004, 1051, 1071, 1144, 1306, 1363, 1398, 1491, 1602, 1630, 1663, 1770, 3080, 3166, 3474, 3526, 3648
Li ⁺ (C ₁)	0.17 (0.02)	126, 175, 203, 236, 339, 373, 398, 472, 531, 543, 567, 596, 644, 682, 730, 749, 778, 906, 949, 976, 1105, 1122, 1223, 1320, 1372, 1437, 1526, 1550, 1636, 1678, 1690, 3164, 3177, 3474, 3489, 3612
Li ⁺ (C ₂)	0.19 (0.02)	91, 164, 195, 238, 264, 381, 425, 443, 448, 477, 515, 547, 592, 644, 722, 762, 788, 942, 976, 977, 1069, 1119, 1208, 1282, 1340, 1409, 1432, 1524, 1590, 1633, 1680, 3146, 3161, 3500, 3571, 3622
Li ⁺ (C ₃)	0.19 (0.02)	111, 192, 227, 260, 311, 360, 420, 430, 435, 499, 538, 572, 589, 656, 726, 774, 787, 927, 971, 984, 1067, 1126, 1201, 1291, 1334, 1404, 1466, 1541, 1575, 1637, 1677, 3148, 3162, 3486, 3581, 3613
Li ⁺ (C ₄)	0.19 (0.02)	73, 144, 147, 182, 372, 381, 462, 512, 553, 599, 604, 612, 649, 696, 713, 757, 779, 913, 964, 984, 1105, 1215, 1224, 1294, 1369, 1392, 1462, 1537, 1610, 1683, 1824, 3159, 3176, 3392, 3484, 3492
Li ⁺ (C ₅)	0.19 (0.02)	81, 95, 120, 188, 352, 370, 463, 500, 513, 521, 602, 635, 651, 689, 756, 784, 838, 889, 962, 996, 1045, 1114, 1204, 1272, 1373, 1392, 1442, 1509, 1641, 1702, 1716, 3174, 3189, 3371, 3475, 3517
Li ⁺ (C ₆)	0.18 (0.02)	128, 182, 210, 250, 253, 358, 415, 436, 461, 523, 571, 603, 627, 675, 728, 754, 790, 951, 955, 1003, 1056, 1134, 1235, 1326, 1350, 1405, 1530, 1597, 1625, 1685, 1689, 3135, 3175, 3443, 3485, 3601

Table 1S (continued). Vibrational Frequencies and Average Vibrational Energies of Cytosine and M⁺(cytosine)^a

Species	E _{int} , eV	Vibrational Frequencies ^a , cm ⁻¹
Na ⁺ (C ₁)	0.20 (0.02)	74, 125, 160, 197, 263, 271, 373, 394, 518, 535, 546, 573, 641, 680, 727, 742, 775, 905, 941, 975, 1101, 1120, 1222, 1300, 1367, 1434, 1519, 1554, 1633, 1682, 1696, 3162, 3177, 3483, 3498, 3622
Na ⁺ (C ₂)	0.20 (0.02)	59, 147, 187, 194, 236, 243, 360, 431, 480, 506, 511, 542, 584, 682, 762, 764, 804, 966, 968, 972, 1060, 1104, 1202, 1263, 1311, 1384, 1406, 1497, 1557, 1610, 1641, 3106, 3125, 3486, 3570, 3595
Na ⁺ (C ₃)	0.20 (0.02)	65, 145, 184, 213, 237, 241, 348, 428, 476, 498, 525, 552, 585, 647, 714, 771, 775, 916, 973, 978, 1068, 1122, 1218, 1291, 1336, 1407, 1455, 1532, 1579, 1628, 1670, 3138, 3159, 3492, 3575, 3620
Na ⁺ (C ₄)	0.20 (0.02)	48, 81, 148, 179, 242, 391, 395, 510, 547, 551, 582, 600, 691, 706, 726, 760, 778, 944, 949, 968, 1086, 1184, 1199, 1265, 1356, 1375, 1441, 1498, 1585, 1643, 1807, 3128, 3143, 3364, 3482, 3488
Na ⁺ (C ₅)	0.20 (0.02)	54, 59, 120, 177, 249, 358, 368, 499, 507, 515, 574, 594, 632, 694, 754, 769, 836, 887, 958, 988, 1048, 1115, 1202, 1277, 1373, 1394, 1444, 1486, 1642, 1711, 1720, 3172, 3189, 3366, 3482, 3525
Na ⁺ (C ₆)	0.20 (0.02)	76, 145, 167, 177, 203, 243, 361, 416, 424, 517, 546, 596, 624, 672, 724, 748, 787, 943, 952, 1006, 1057, 1126, 1211, 1325, 1351, 1403, 1523, 1603, 1623, 1681, 1700, 3125, 3175, 3450, 3492, 3609
K ⁺ (C ₁)	0.21 (0.02)	50, 86, 154, 173, 200, 222, 367, 390, 511, 534, 537, 568, 642, 682, 726, 740, 773, 903, 936, 973, 1096, 1119, 1220, 1289, 1363, 1431, 1513, 1558, 1630, 1681, 1712, 3160, 3175, 3488, 3504, 3629
K ⁺ (C ₂)	0.19 (0.02)	-51 , 8, 171, 191, 225, 353, 430, 472, 503, 542, 546, 581, 685, 765, 766, 801, 963, 966, 970, 1060, 1104, 1211, 1265, 1310, 1381, 1412, 1494, 1558, 1607, 1636, 3101, 3122, 3490, 3566, 3601
K ⁺ (C ₃)	0.21 (0.02)	50, 112, 148, 164, 186, 225, 345, 430, 496, 509, 521, 548, 585, 647, 717, 769, 772, 919, 974, 976, 1069, 1119, 1229, 1291, 1338, 1407, 1446, 1528, 1583, 1624, 1667, 3130, 3158, 3497, 3574, 3626
K ⁺ (C ₄)	0.20 (0.02)	47(2), 115, 175, 175, 362, 363, 500, 503, 520, 570, 625, 662, 698, 737, 763, 840, 880, 965, 980, 1063, 1123, 1211, 1311, 1374, 1396, 1451, 1479, 1630, 1707, 1738, 3160, 3175, 3406, 3481, 3531
K ⁺ (C ₅)	0.21 (0.02)	47, 51, 120, 171, 173, 359, 366, 499(2), 517, 563, 588, 631, 701, 753, 766, 835, 886, 958, 983, 1050, 1117, 1201, 1279, 1374, 1396, 1445, 1472, 1641, 1711, 1732, 3169, 3188, 3360, 3484, 3530
K ⁺ (C ₆)	0.22 (0.02)	34, 64, 108, 166, 180, 197, 353, 409, 422, 512, 540, 592, 621, 670, 721, 745, 786, 937, 954, 1008, 1057, 1118, 1193, 1323, 1353, 1401, 1518, 1609, 1622, 1678, 1715, 3115, 3174, 3454, 3497, 3615

^aObtained from vibrational analyses of the optimized geometries as described in the text and scaled by 0.9646. The imaginary frequencies (indicated in boldface) correspond to the out-of-plane vibration(s) of the amino group.

Table 2S. Vibrational Frequencies (in cm^{-1}) of the Transition States for the Unimolecular Tautomerization of $\text{M}^+(\text{Cytosine})^a$

Species	E_{int} , eV	Vibrational Frequencies ^a , cm^{-1}
$\text{Li}^+(\text{C})\text{-TS}_{1,2}$	0.18 (0.02)	-1832 , 108, 191, 194, 205, 323, 374, 408, 454, 513, 552, 571, 631, 642, 694, 763, 810, 924, 978, 982, 999, 1098, 1151, 1308, 1330, 1414, 1486, 1530, 1586, 1627, 1670, 1980, 3158, 3170, 3488, 3611
$\text{Li}^+(\text{C})\text{-TS}_{2,3}$	0.17 (0.02)	-137 , 85, 178, 223, 318, 323, 434, 450, 520, 534, 541, 557, 591, 660, 727, 756, 800, 938, 965, 970, 1063, 1112, 1225, 1248, 1323, 1379, 1450, 1524, 1557, 1619, 1667, 3143, 3158, 3487, 3541, 3614
$\text{Li}^+(\text{C})\text{-TS}_{3,6}$	0.18 (0.02)	-1827 , 110, 200, 212, 218, 348, 350, 418, 431, 516, 545, 602, 622, 645, 688, 762, 778, 932, 976, 988, 988, 1131, 1151, 1282, 1349, 1419, 1483, 1536, 1601, 1621, 1684, 1986, 3149, 3164, 3481, 3603
$\text{Li}^+(\text{C})\text{-TS}_{1,4}$	0.18 (0.02)	-1853 , 80, 87, 132, 204, 378, 382, 468, 547, 564, 583, 615, 643, 662, 698, 751, 816, 902, 971, 1026, 1071, 1076, 1132, 1226, 1373, 1376, 1439, 1470, 1604, 1691, 1702, 1986, 3163, 3184, 3482, 3501
$\text{Li}^+(\text{C})\text{-TS}_{4,5}$	0.16 (0.02)	-993 , 114, 130, 156, 251, 393, 444, 501, 549, 557, 692, 703, 730, 767, 802, 811, 930, 955, 1017, 1063, 1071, 1145, 1212, 1319, 1472, 1486, 1533, 1572, 1768, 1796, 1979, 3299, 3322, 3646, 3677, 3902
$\text{Na}^+(\text{C})\text{-TS}_{1,2}$	0.17 (0.02)	-1826 , 76, 101, 211, 222, 259, 397, 405, 453, 571, 588, 610, 688, 726, 824, 865, 869, 1047, 1054, 1097, 1126, 1178, 1216, 1336, 1397, 1501, 1582, 1638, 1691, 1745, 1769, 2168, 3297, 3314, 3701, 3819
$\text{Na}^+(\text{C})\text{-TS}_{2,3}$	0.18 (0.02)	-83 , 54, 173, 202, 290, 331, 356, 435, 468, 518, 540, 544, 590, 662, 735, 762, 796, 938, 968, 974, 1071, 1115, 1229, 1265, 1325, 1386, 1446, 1519, 1565, 1618, 1664, 3139, 3156, 3484, 3554, 3609
$\text{Na}^+(\text{C})\text{-TS}_{3,6}$	0.19 (0.02)	-1820 , 68, 132, 194, 201, 230, 297, 351, 420, 505, 537, 597, 623, 645, 689, 757, 777, 930, 974, 987, 1005, 1127, 1153, 1282, 1346, 1403, 1488, 1530, 1610, 1617, 1681, 2010, 3142, 3164, 3488, 3610
$\text{Na}^+(\text{C})\text{-TS}_{1,4}$	0.19 (0.02)	-1857 , 44, 57, 129, 195, 264, 374, 386, 524, 555, 566, 582, 633, 642, 698, 745, 784, 902, 968, 1023, 1063, 1079, 1130, 1224, 1374, 1379, 1421, 1470, 1605, 1686, 1715, 1991, 3160, 3184, 3485, 3506
$\text{Na}^+(\text{C})\text{-TS}_{4,5}$	0.18 (0.02)	-1011 , 74(2), 158, 237, 299, 399, 445, 545, 555, 619, 677, 691, 761, 782, 797, 927, 951, 1001, 1058, 1076, 1145, 1211, 1319, 1477, 1484, 1524, 1560, 1780, 1794, 1971, 3294, 3320, 3655, 3686, 3915
$\text{K}^+(\text{C})\text{-TS}_{1,2}$	0.20 (0.02)	-1821 , 47, 80, 164, 184, 188, 214, 373, 405, 496, 548, 554, 636, 642, 699, 755, 816, 923, 971, 981, 1030, 1094, 1150, 1297, 1326, 1401, 1486, 1532, 1587, 1628, 1658, 2021, 3155, 3168, 3503, 3628
$\text{K}^+(\text{C})\text{-TS}_{2,3}$	0.18 (0.02)	-51 , 53, 167, 178, 237, 331, 384, 435, 469, 512, 527, 543, 590, 659, 730, 765, 791, 934, 970, 979, 1077, 1117, 1234, 1276, 1326, 1391, 1447, 1516, 1571, 1618, 1661, 3133, 3155, 3482, 3568, 3606
$\text{K}^+(\text{C})\text{-TS}_{3,6}$	0.20 (0.02)	-1814 , 53, 104, 169, 188, 197, 248, 347, 423, 497, 534, 591, 623, 647, 691, 753, 777, 934, 971, 987, 1018, 1124, 1153, 1281, 1341, 1392, 1494, 1527, 1612, 1621, 1679, 2026, 3134, 3163, 3492, 3616
$\text{K}^+(\text{C})\text{-TS}_{1,4}$	0.20 (0.02)	-1861 , 35, 49, 127, 188, 189, 370, 386, 520, 538, 562, 579, 631, 641, 702, 740, 778, 902, 967, 1022, 1057, 1081, 1128, 1222, 1373, 1381, 1408, 1473, 1605, 1683, 1729, 1994, 3157, 3184, 3487, 3509

Table 2S. (continued) Vibrational Frequencies (in cm^{-1}) of the Transition States for the Unimolecular Tautomerization of $\text{M}^+(\text{Cytosine})$ ^a

Species	E_{int} , eV	Vibrational Frequencies ^a , cm^{-1}
$\text{K}^+(\text{C})\text{-TS}_{4,5}$	0.18 (0.02)	-1029 , 58(2), 159, 177, 228, 399, 444, 546, 554, 597, 662, 681, 754, 779, 794, 924, 947, 997, 1053, 1081, 1143, 1212, 1317, 1476, 1482, 1515, 1556, 1790, 1797, 1964, 3291, 3321, 3661, 3692, 3923

^aObtained from a vibrational analysis of the MP2(full)/6-31G* geometry optimized structures and scaled by 0.9646. The imaginary frequencies (indicated in boldface) correspond to the reaction coordinate for the tautomerization.

Table 3S. Rotational Constants of $\text{M}^+(\text{cytosine})$ in cm^{-1}

Reactant	Energized Molecule		Transition State		
	1-D ^a	2-D ^b	1-D ^a	2-D ^c	2-D ^{b,d}
$\text{Li}^+(\text{C}_1)$	0.097	0.049	0.131	0.055	0.060
$\text{Li}^+(\text{C}_2)$	0.097	0.050	0.131	0.054	0.060
$\text{Li}^+(\text{C}_3)$	0.128	0.044	0.129	0.055	0.064
$\text{Li}^+(\text{C}_4)$	0.114	0.041	0.128	0.054	0.068
$\text{Li}^+(\text{C}_5)$	0.117	0.040	0.129	0.054	
$\text{Li}^+(\text{C}_6)$	0.130	0.044	0.130	0.054	
$\text{Na}^+(\text{C}_1)$	0.077	0.036	0.131	0.055	0.0070
$\text{Na}^+(\text{C}_2)$	0.075	0.037	0.131	0.054	0.0063
$\text{Na}^+(\text{C}_3)$	0.124	0.029	0.129	0.055	0.0085
$\text{Na}^+(\text{C}_4)$	0.106	0.024	0.128	0.054	0.010
$\text{Na}^+(\text{C}_5)$	0.111	0.024	0.129	0.054	
$\text{Na}^+(\text{C}_6)$	0.111	0.024	0.130	0.054	
$\text{K}^+(\text{C}_1)$	0.074	0.025	0.131	0.055	0.0036
$\text{K}^+(\text{C}_2)$	0.071	0.026	0.131	0.054	0.0036
$\text{K}^+(\text{C}_3)$	0.123	0.021	0.129	0.055	0.0043
$\text{K}^+(\text{C}_4)$	0.104	0.017	0.128	0.054	0.0054
$\text{K}^+(\text{C}_5)$	0.109	0.016	0.129	0.054	
$\text{K}^+(\text{C}_6)$	0.127	0.021	0.130	0.054	

^aActive external. ^bInactive external. ^cRotational constants of the transition state treated as free internal rotors.
^dTwo-dimensional rotational constant of the transition state at the threshold energy for dissociation, treated variationally and statistically.

Table 4S. Rotational Constants for Unimolecular Tautomerization of M^+ (cytosine) Complexes in cm^{-1}

Process	Energized Molecule		Transition State	
	1-D ^a	2-D ^b	1-D ^a	2-D ^b
$\text{Li}^+(\text{C}_2) \rightarrow \text{Li}^+(\text{C}_3)$	0.097	0.050	0.128	0.044
$\text{Na}^+(\text{C}_2) \rightarrow \text{Na}^+(\text{C}_3)$	0.075	0.036	0.085	0.035
$\text{K}^+(\text{C}_2) \rightarrow \text{K}^+(\text{C}_3)$	0.072	0.026	0.073	0.027
$\text{Li}^+(\text{C}_2) \rightarrow \text{Li}^+(\text{C}_1)$	0.097	0.050	0.097	0.049
$\text{Na}^+(\text{C}_2) \rightarrow \text{Na}^+(\text{C}_1)$	0.075	0.036	0.077	0.036
^a Active external. ^b Inactive external.				

Table 5S. MP2(full)/6-31G* Optimized Geometries of cytosine and M⁺(cytosine)^a

C ₁			C ₂			C ₃		
x, y, z			x, y, z			x, y, z		
N	0.082054,	-1.025299, -0.000345	N	0.048633,	-1.005303, -0.000279	N	-0.006719,	-0.977298, -0.000008
C	-1.190120,	-0.521040, 0.000025	C	-1.104881,	-0.335882, -0.000697	C	1.134373,	-0.274980, -0.000019
N	-1.277603,	0.891306, -0.000460	N	-1.303504,	0.986104, -0.000173	N	1.296524,	1.047206, -0.000004
C	-0.188685,	1.696894, -0.000143	C	-0.170585,	1.718193, 0.000109	C	0.138854,	1.736292, 0.000004
C	1.0452331,	0.180541, 0.000522	C	1.093093,	1.162525, 0.000110	C	-1.113217,	1.148959, 0.000003
C	1.129889,	-0.260218, -0.000021	C	1.156166,	-0.246405, 0.000058	C	-1.141633,	-0.257701, 0.000001
H	-2.202156,	1.268127, -0.000665	H	-0.304054,	2.798396, 0.000241	H	0.237684,	2.820567, 0.000008
H	-0.369506,	2.751702, 0.000072	H	1.985912,	1.778956, 0.000220	H	-2.021554,	1.742414, 0.000006
H	1.917553,	1.794551, 0.001168	N	2.337432,	-0.919227, 0.000104	N	-2.302873,	-0.966464, 0.000002
N	2.339955,	-0.844685, 0.000257	H	2.313351,	-1.928373, 0.000029	H	-2.254876,	-1.974322, 0.000000
H	2.371902,	-1.841570, -0.000010	H	3.224787,	-0.442785, 0.000229	H	-3.202718,	-0.513797, 0.000006
H	3.188205,	-0.325928, 0.000677	O	-2.210038,	-1.117123, 0.000427	O	2.278575,	-0.997758, 0.000012
O	-2.211843,	-1.171650, 0.000488	H	-2.960380,	-0.490815, 0.000823	H	1.994079,	-1.932313, 0.000020

C ₄			C ₅			C ₆		
x, y, z			x, y, z			x, y, z		
N	-0.034986,	-0.975596, -0.000141	N	0.049275,	-0.973002, -0.000031	N	-0.026097,	-0.907268, -0.000011
C	1.226164,	-0.421340, -0.000603	C	-1.224301,	-0.424771, -0.000099	C	1.296613,	-0.374482, -0.000032
N	1.206793,	0.970524, -0.000020	N	-1.220193,	0.959335, -0.000007	N	1.394982,	1.001886, -0.000001
C	0.047526,	1.716155, 0.000072	C	-0.064325,	1.721278, 0.000014	C	0.274888,	1.700144, 0.000005
C	-1.169647,	1.130575, -0.000038	C	1.156100,	1.148088, -0.000008	C	-1.037662,	1.197243, 0.000000
C	-1.277214,	-0.322145, 0.000054	C	1.287848,	-0.300037, -0.000003	C	-1.169163,	-0.176820, -0.000001
H	0.182326,	2.793052, 0.000290	H	-0.212950,	2.795847, 0.000060	H	0.406878,	2.783088, 0.000017
H	-2.070072,	1.733359, 0.000043	H	2.063656,	1.737178, 0.000014	H	-1.901424,	1.850022, 0.000008
H	-2.327367,	-1.070700, 0.000110	N	2.447266,	-0.862960, 0.000026	N	-2.351598,	-0.847266, 0.000007
N	-3.164983,	-0.483227, 0.000032	H	2.364426,	-1.885957, 0.000030	H	-2.397503,	-1.854435, -0.000001
O	2.266388,	-1.068915, 0.000335	O	-2.249031,	-1.097352, 0.000061	H	-3.220606,	-0.336626, 0.000011
H	2.118441,	1.410524, 0.000265	H	-2.136336,	1.389770, 0.000050	O	2.231706,	-1.165828, 0.000023
H	-0.068873,	-1.991447, 0.000142	H	0.047087,	-1.988974, 0.000011	H	-0.050049,	-1.923405, -0.000008

Li ⁺ (C ₁)			Li ⁺ (C ₂)			Li ⁺ (C ₃)		
x, y, z			x, y, z			x, y, z		
N	-0.050600,	-0.907629, 0.000013	N	-0.058955,	-0.866651, -0.00012	N	-0.250167,	-1.023062, -0.00009
C	-1.169617,	-0.123258, 0.000035	C	-1.069276,	0.019086, 0.000013	C	0.906143,	-0.404814, -0.000103
N	-1.028158,	1.241203, 0.000007	N	-1.059954,	1.322168, 0.000141	N	1.194166,	0.896039, 0.000002
C	0.198377,	1.840496, 0.000011	C	0.194859,	1.852331, 0.000159	C	0.100135,	1.704816, 0.000076
C	1.323954,	1.082327, 0.000005	C	1.331564,	1.081216, 0.000050	C	-1.177165,	1.202732, 0.000068
C	1.163108,	-0.339838, -0.000004	C	1.181370,	-0.326677, -0.000114	C	-1.333014,	-0.206867, -0.00001
H	-1.886724,	1.787610, -0.000011	H	0.249752,	2.936880, 0.000310	H	0.290345,	2.773492, 0.000087
H	0.208147,	2.924807, 0.000001	H	2.316005,	1.535828, 0.000141	H	-2.034767,	1.866167, 0.000088
H	2.304153,	1.542867, -0.000002	N	2.235857,	-1.156941, -0.000061	N	-2.529812,	-0.802414, -0.00004
N	2.233894,	-1.138382, -0.000019	H	2.110368,	-2.159614, -0.000575	H	-2.581158,	-1.813706, -0.00009
H	2.122066,	-2.144231, -0.000027	H	3.181883,	-0.800773, -0.000224	H	-3.391739,	-0.274235, -0.00006
H	3.174249,	-0.766001, -0.000032	O	-2.279020,	-0.648141, -0.00045	O	2.062603,	-1.156066, 0.000191
O	-2.306329,	-0.659162, -0.00006	H	-3.009338,	0.005490, -0.000419	H	1.842012,	-2.110094, 0.000290
Li	-1.550713,	-2.398823, 0.000092	Li	-1.422086,	-2.392815, 0.001362	Li	3.166193,	0.512585, -0.000395

^aStandard Orientation, Å.

Table S5 (continued). MP2(full)/6-31G* Optimized Geometries of Cytosine and M⁺(cytosine)^a

Li ⁺ (C ₁)			Li ⁺ (C ₂)			Li ⁺ (C ₃)		
x, y, z			x, y, z			x, y, z		
N	-0.050600, -0.907629, 0.000013		N	-0.058955, -0.866651, -0.00012		N	-0.250167, -1.023062, -0.00009	
C	-1.169617, -0.123258, 0.000035		C	-1.069276, 0.019086, 0.000013		C	0.906143, -0.404814, -0.000103	
N	-1.028158, 1.241203, 0.000007		N	-1.059954, 1.322168, 0.000141		N	1.194166, 0.896039, 0.000002	
C	0.198377, 1.840496, 0.000011		C	0.194859, 1.852331, 0.000159		C	0.100135, 1.704816, 0.000076	
C	1.323954, 1.082327, 0.000005		C	1.331564, 1.081216, 0.000050		C	-1.177165, 1.202732, 0.000068	
C	1.163108, -0.339838, -0.000004		C	1.181370, -0.326677, -0.000114		C	-1.333014, -0.206867, -0.000001	
H	-1.886724, 1.787610, -0.000011		H	0.249752, 2.936880, 0.000310		H	0.290345, 2.773492, 0.000087	
H	0.208147, 2.924807, 0.000001		H	2.316005, 1.535828, 0.000141		H	-2.034767, 1.866167, 0.000088	
H	2.304153, 1.542867, -0.000002		N	2.235857, -1.156941, -0.000061		N	-2.529812, -0.802414, -0.000004	
N	2.233894, -1.138382, -0.000019		H	2.110368, -2.159614, -0.000575		H	-2.581158, -1.813706, -0.000009	
H	2.122066, -2.144231, -0.000027		H	3.181883, -0.800773, -0.000224		H	-3.391739, -0.274235, -0.000006	
H	3.174249, -0.766001, -0.000032		O	-2.279020, -0.648141, -0.000045		O	2.062603, -1.156066, 0.000191	
O	-2.306329, -0.659162, -0.000006		H	-3.009338, 0.005490, -0.000419		H	1.842012, -2.110094, 0.000290	
Li	-1.550713, -2.398823, 0.000092		Li	-1.422086, -2.392815, 0.001362		Li	3.166193, 0.512585, -0.000395	

Li ⁺ (C ₄)			Li ⁺ (C ₅)			Li ⁺ (C ₆)		
x, y, z			x, y, z			x, y, z		
N	-0.015283, -0.886334, -0.000002		N	-0.002646, -0.900579, -0.000004		N	0.229790, -0.948051, -0.000554	
C	-1.053343, -0.036154, -0.000009		C	1.052658, -0.056956, -0.000011		C	-1.065009, -0.449860, -0.001156	
N	-0.743820, 1.283093, -0.000002		N	0.766821, 1.260451, -0.000001		C	1.339211, -0.158669, 0.000060	
C	0.572331, 1.741917, 0.000003		C	-0.545119, 1.747481, 0.000001		H	0.299618, -1.965466, -0.000559	
C	1.607135, 0.880814, 0.000004		C	-1.589545, 0.902229, -0.000004		N	-1.250669, 0.894427, -0.000879	
C	1.369532, -0.554792, 0.000004		C	-1.388562, -0.536711, -0.000001		C	1.135290, 1.226709, 0.000652	
H	-1.519758, 1.935616, -0.000006		H	1.553645, 1.899958, 0.000015		N	2.541523, -0.743473, -0.000025	
H	0.688306, 2.819301, 0.000005		H	-0.638845, 2.826534, 0.000006		O	-2.043209, -1.237040, 0.000683	
H	2.625070, 1.252133, 0.000006		H	-2.610268, 1.262928, -0.000002		C	-0.167395, 1.686765, -0.000036	
N	2.166140, -1.554796, 0.000000		N	-2.368722, -1.356114, 0.000003		H	1.975085, 1.910130, 0.001329	
H	3.142904, -1.252019, -0.000008		H	-2.084146, -2.340634, 0.000005		H	2.664108, -1.748303, -0.000154	
O	-2.255712, -0.427423, -0.000006		O	2.244998, -0.480377, 0.000016		H	3.381169, -0.178369, 0.000099	
H	-0.212968, -1.885427, -0.000003		H	0.231568, -1.890335, 0.000009		H	-0.362740, 2.756077, -0.000201	
Li	-3.833687, -1.178224, 0.000023		Li	3.881103, -1.092666, -0.000019		Li	-3.236217, 0.290752, 0.000370	

Na ⁺ (C ₁)			Na ⁺ (C ₂)			Na ⁺ (C ₃)		
x, y, z			x, y, z			x, y, z		
N	-0.259786, 0.540524, 0.000081		N	0.243011, 0.502429, -0.000008		N	-0.775696, -1.024459, -0.000003	
C	-0.609842, -0.784103, -0.000002		C	0.491375, -0.813755, -0.000061		C	0.443314, -0.519508, -0.000027	
N	0.397692, -1.728696, -0.000048		N	-0.337467, -1.828210, -0.000003		N	0.846330, 0.751044, 0.000010	
C	1.718009, -1.395257, -0.000104		C	-1.645647, -1.475376, 0.000009		C	-0.175836, 1.650094, 0.000035	
C	2.083684, -0.088652, -0.000071		C	-2.067857, -0.165009, 0.000000		C	-1.497717, 1.272622, 0.000019	
C	0.033676, 0.883071, 0.000050		C	-1.070868, 0.842031, 0.000018		C	-1.779810, -0.116428, -0.000001	
H	0.097299, -2.700170, -0.000011		H	-2.359402, -2.294227, -0.000002		H	0.108551, 2.698554, 0.000048	
H	2.427466, -2.215437, -0.000151		H	-3.120791, 0.091424, -0.000013		H	-2.289750, 2.013623, 0.000024	
H	3.127464, 0.199481, -0.000112		N	-1.385700, 2.151727, 0.000019		N	-3.029752, -0.603444, -0.000002	
N	1.338028, 2.188040, 0.000065		H	-0.667870, 2.860316, 0.000011		H	-3.170451, -1.605732, -0.000004	
H	0.602606, 2.881751, 0.000226		H	-2.347664, 2.456543, 0.000007		H	-3.839861, 0.000452, -0.000012	
H	2.295837, 2.510582, 0.000099		O	1.835033, -1.092374, 0.000060		O	1.494998, -1.408648, 0.000069	
O	-1.804871, -1.150411, 0.000277		H	1.935826, -2.062805, 0.000100		H	1.121832, -2.314490, 0.000104	
Na	-2.708764, 0.894332, -0.000188		Na	0.545337, 1.052616, -0.000022		Na	3.171256, 0.263101, -0.000046	

Table S5 (continued). MP2(full)/6-31G* Optimized Geometries of Cytosine and M⁺(cytosine)^a

Na ⁺ (C ₄)			Na ⁺ (C ₅)			Na ⁺ (C ₆)		
x, y, z			x, y, z			x, y, z		
N	0.333700, -0.800135, -0.000024		N	-0.372955, -0.827594, 0.000022		N	-0.718857, -0.947240, 0.000727	
C	-0.530924, 0.236966, -0.000005		C	0.534743, 0.182332, 0.000033		C	0.613825, -0.533374, -0.008496	
N	0.044054, 1.473337, -0.000001		N	0.015740, 1.433894, 0.000011		N	0.869771, 0.805147, -0.008111	
C	1.423927, 1.662927, 0.000005		C	-1.358267, 1.686051, -0.000018		C	-0.173372, 1.646412, -0.005102	
C	2.271789, 0.614893, -0.000002		C	-2.243726, 0.675211, -0.000029		C	-1.504194, 1.266877, -0.001123	
C	1.754025, -0.746430, 0.000005		C	-1.796307, -0.706702, -0.000007		C	-1.783511, -0.101946, 0.002247	
H	-0.587021, 2.266127, 0.000026		H	0.679333, 2.199467, 0.000032		H	0.077175, 2.704797, -0.005791	
H	1.750873, 2.696557, 0.000002		H	-1.637884, 2.732651, -0.000031		H	-2.301921, 1.998755, 0.000370	
H	3.343074, 0.778661, -0.000005		H	-3.311023, 0.855287, -0.000052		N	-3.017549, -0.623860, 0.007966	
N	2.344974, -1.883595, 0.000014		N	-2.623217, -1.682753, -0.000002		H	-3.191242, -1.620480, 0.009341	
H	3.361273, -1.768112, 0.000010		H	-2.168027, -2.601178, 0.000003		H	-3.825729, -0.015696, 0.009771	
O	-1.779232, 0.088987, -0.000001		O	1.773188, -0.030046, 0.000071		O	1.515854, -1.396842, -0.011464	
H	-0.055559, -1.740417, -0.000002		H	0.025697, -1.762587, 0.000049		H	-0.846231, -1.958383, 0.001534	
Na	-3.831889, -0.462009, 0.000006		Na	3.842797, -0.424496, -0.000052		Na	3.191916, 0.159964, 0.013386	

K ⁺ (C ₁)			K ⁺ (C ₂)			K ⁺ (C ₃)		
x, y, z			x, y, z			x, y, z		
N	0.033315, 0.429114, 0.000139		N	-0.027527, 0.363210, -0.000039		N	1.234418, -1.015598, -0.000068	
C	-0.056088, -0.939275, -0.000099		C	-0.120877, -0.970604, 0.000035		C	-0.006001, -0.553732, -0.000007	
N	1.124124, -1.668894, -0.000101		N	-1.188315, -1.740573, 0.000026		N	-0.449863, 0.702916, 0.000038	
C	2.354553, -1.089499, -0.000204		C	-2.361395, -1.068989, -0.000004		C	0.546193, 1.628792, 0.000091	
C	2.459575, 0.263239, -0.000164		C	-2.436431, 0.306499, 0.000020		C	1.881946, 1.299689, 0.000045	
C	1.237298, 1.007045, 0.000069		C	-1.213424, 1.019889, -0.000043		C	2.207940, -0.077804, -0.000026	
H	1.016386, -2.679504, -0.000014		H	-3.260335, -1.679292, 0.000107		H	0.230806, 2.668958, 0.000124	
H	3.209160, -1.757280, -0.000277		H	-3.388762, 0.823973, 0.000121		H	2.648388, 2.066997, 0.000052	
H	3.426693, 0.750289, -0.000228		N	-1.185054, 2.370036, 0.000005		N	3.475274, -0.525566, -0.000065	
N	1.286805, 2.349692, 0.000180		H	-0.309468, 2.869468, -0.000042		H	3.647019, -1.522437, -0.000126	
H	0.430926, 2.886541, 0.000355		H	-2.037177, 2.909720, 0.000046		H	4.264952, 0.103939, -0.000068	
H	2.164252, 2.850143, 0.000122		O	1.091975, -1.605116, -0.000029		O	-1.010427, -1.490409, 0.000187	
O	-1.145065, -1.543116, 0.000566		H	0.899508, -2.561683, -0.000023		H	-0.574763, -2.367327, 0.000245	
K	-2.850981, 0.372437, -0.000191		K	2.787341, 0.411014, 0.000014		K	-3.143803, 0.161010, -0.000068	

K ⁺ (C ₄)			K ⁺ (C ₅)			K ⁺ (C ₆)		
x, y, z			x, y, z			x, y, z		
N	-0.773647, -0.774165, -0.000003		N	0.826830, -0.808417, -0.000062		N	1.168808, -0.940686, -0.000064	
C	0.036611, 0.310697, 0.000004		C	-0.039121, 0.244337, -0.000111		C	-0.179363, -0.557080, 0.000337	
N	-0.617993, 1.511324, 0.000002		N	0.549663, 1.470615, -0.000028		N	-0.459370, 0.779258, 0.000557	
C	-2.003234, 1.616991, -0.000001		C	1.932643, 1.652381, 0.000046		C	0.571766, 1.631786, 0.000384	
C	-2.787908, 0.522299, -0.000003		C	2.768098, 0.599420, 0.000061		C	1.913407, 1.282465, 0.000018	
C	-2.190049, -0.803782, -0.000001		C	2.251486, -0.757849, 0.000015		C	2.217860, -0.078519, -0.000190	
H	-0.036258, 2.340253, 0.000004		H	-0.073809, 2.268639, -0.000029		H	0.305339, 2.687024, 0.000504	
H	-2.392787, 2.628610, 0.000002		H	2.265618, 2.683458, 0.000094		H	2.695200, 2.031293, -0.000091	
H	-3.866633, 0.622428, 0.000003		H	3.842931, 0.726477, 0.000120		N	3.462871, -0.580469, -0.000521	
N	-2.717520, -1.971527, 0.000004		N	3.030226, -1.774541, 0.000044		H	3.651767, -1.573786, -0.000613	
H	-3.738074, -1.910120, 0.000023		H	2.525735, -2.667308, 0.000007		H	4.260073, 0.041156, -0.000618	
O	1.283735, 0.234282, 0.000004		O	-1.280368, 0.093351, -0.000119		O	-1.048291, -1.447389, 0.000849	
H	-0.330678, -1.689559, -0.000004		H	0.383539, -1.722753, -0.000087		H	1.314055, -1.948917, -0.000083	
K	3.711903, -0.268554, -0.000001		K	-3.738250, -0.246245, 0.000058		K	-3.167804, 0.098090, -0.000473	

Table 6S. MP2(full)/6-31G* Optimized Geometries of Transition States for Unimolecular Tautomerization of Cytosine and M⁺(cytosine) Complexes.^a

C-TS _{1,2}			C-TS _{2,3}			C-TS _{3,6}		
x, y, z			x, y, z			x, y, z		
N	-0.136188,	1.088360, 0.000176	N	-0.830699,	-0.048605, -0.519890	N	0.004375,	-0.901962, 0.002031
C	1.075642,	-0.521443, -0.001252	C	-0.822213,	-0.096984, 0.814539	C	-1.230879,	-0.285446, 0.001416
N	1.308951,	0.830533, 0.000823	N	0.232684,	-0.043552, 1.625709	N	-1.414997,	1.039964, 0.006518
C	0.281816,	1.691596, 0.002443	C	1.419976,	0.069063, 0.998712	C	-0.260860,	1.730463, 0.003176
C	-1.001645,	1.192915, -0.001383	C	1.552285,	0.126590, -0.375101	C	1.021431,	1.192347, -0.004276
C	-1.157334,	-0.218644, -0.001445	C	0.364093,	0.064861, -1.119681	C	1.144351,	-0.205033, -0.001621
H	0.506617,	2.754187, 0.004309	H	2.295254,	0.113988, 1.643806	H	-0.373867,	2.813642, 0.006226
H	-1.859362,	1.855058, -0.013141	H	2.523448,	0.223203, -0.849720	H	1.896671,	1.831412, -0.013474
N	-2.402038,	-0.772658, -0.056609	N	0.346786,	0.181017, -2.489314	N	2.332488,	-0.875019, -0.064882
H	-2.441306,	-1.766883, 0.131094	H	-0.524313,	-0.115487, -2.913993	H	2.313416,	-1.852729, 0.196524
H	-3.197690,	-0.225444, 0.239386	H	1.178441,	-0.108320, -2.987463	H	3.159131,	-0.371975, 0.228659
O	2.240969,	-1.070330, 0.003961	O	-2.044651,	-0.287200, 1.411022	O	-2.083329,	-1.248985, 0.001438
H	2.478051,	0.292569, 0.005763	H	-2.441870,	0.581016, 1.592843	H	-0.826044,	-1.863341, -0.004269

C-TS _{1,4}			C-TS _{4,5}			Li ⁺ (C)-TS _{1,2}		
x, y, z			x, y, z			x, y, z		
N	0.000000,	0.974535, 0.000000	N	-0.057780,	-0.963843, 0.034987	N	0.018503	-0.962177 -0.001065
C	1.221963,	0.372137, 0.000000	C	1.222130,	-0.453978, 0.002282	C	-1.037092	-0.157415 -0.000644
N	1.116360,	-1.047007, 0.000000	N	1.246723,	0.934183, -0.005550	N	-1.094865	1.171880 -0.000464
C	-0.054195,	-1.740483, 0.000000	C	0.112180,	1.715479, 0.000767	C	0.083405	1.841342 0.000323
C	-1.260206,	-1.109707, 0.000000	C	-1.124838,	1.176102, 0.011698	C	1.256471	1.124035 0.000620
C	-1.199293,	0.317924, 0.000000	C	-1.313910,	-0.276176, 0.011294	C	1.208049	-0.299618 -0.000079
H	2.007437,	-1.530867, 0.000000	H	0.285282,	2.786872, -0.000875	H	0.052496	2.925815 0.000774
H	0.036205,	-2.822718, 0.000000	H	-2.007239,	1.802765, 0.012168	H	2.210802	1.638028 0.001348
N	-2.191603,	-1.660139, 0.000000	N	-2.391390,	-0.894868, -0.064860	N	2.321672	-1.037960 -0.000040
H	-2.075054,	1.299335, 0.000000	H	-3.224465,	-1.376122, 0.178634	H	2.265908	-2.048383 -0.000942
H	-3.080234,	1.157654, 0.000000	O	2.241752,	-1.134749, -0.011813	H	3.239611	-0.613021 0.000412
O	2.323242,	0.905477, 0.000000	H	2.171126,	1.345713, -0.026382	O	-2.288902	-0.588002 -0.000047
H	-0.896501,	1.984985, 0.000000	H	-0.114956,	-1.978102, 0.022684	H	-2.388858	0.743288 0.000556
						Li	-1.616971	-2.397993 0.002632

Li ⁺ (C)-TS _{2,3}			Li ⁺ (C)-TS _{3,6}			Li ⁺ (C)-TS _{1,4}		
x, y, z			x, y, z			x, y, z		
N	-0.090713,	-0.969603, -0.129432	N	0.198203,	-0.946660, -0.000296	N	0.044173,	-0.862863, 0.000000
C	0.963292,	-0.179770, -0.200109	C	-0.997038,	-0.330181, -0.000332	C	1.046324,	0.009312, 0.000000
C	-1.272389,	-0.318032, -0.000693	N	-1.271730,	0.959748, -0.000571	C	-1.301700,	-0.512727, 0.000000
N	1.083355,	1.131407, -0.086448	C	-0.141463,	1.732800, -0.000028	N	0.676004,	1.338234, 0.000001
C	-1.313069,	1.097131, 0.053615	C	1.132310,	1.214981, 0.000205	C	-1.665078,	0.870264, 0.000000
N	-2.371141,	-1.083108, 0.089931	C	1.319087,	-0.197600, 0.000014	N	-1.893377,	-1.672798, 0.000000
O	2.192184,	-0.876689, -0.299451	H	-0.300739,	2.806697, 0.000028	O	2.276975,	-0.285581, -0.000001
Li	2.556146,	-0.714601, 1.546281	H	1.989858,	1.878185, 0.000536	C	-0.635062,	1.755267, 0.000000
C	-0.114543,	1.771378, 0.025313	N	2.511305,	-0.791197, 0.000187	H	1.431215,	2.016843, 0.000001
H	-2.252077,	1.633205, 0.140432	H	2.587197,	-1.801356, 0.000025	H	-2.695056,	1.202999, -0.000001
H	-2.284356,	-2.080778, -0.060626	H	3.366708,	-0.250692, 0.000423	H	-2.903839,	-1.792319, 0.000001
H	-3.297325,	-0.678791, 0.089957	O	-1.899598,	-1.291422, 0.000021	Li	3.605565,	-1.427573, 0.000002
H	2.768366,	-0.345884, -0.891482	H	-0.704311,	-1.934079, 0.000218	H	-0.783171,	2.829670, 0.000000
H	-0.070772,	2.854458, 0.091366	Li	-3.227917,	0.186460, 0.001399	H	-0.536157,	-2.040526, 0.000000

Table 6S (continued). MP2(full)/6-31G* Optimized Geometries of Transition States for Unimolecular Tautomerization of Cytosine and M⁺(cytosine) Complexes.

Na ⁺ (C)-TS _{4,5}			Na ⁺ (C)-TS _{1,2}			Na ⁺ (C)-TS _{2,3}		
x, y, z			x, y, z			x, y, z		
N	0.002373,	-0.887470, 0.000000	N	-0.188781,	0.646303, 0.000014	N	-0.354906,	-0.881098, -0.422653
C	-1.063507,	-0.072173, 0.000000	C	-0.577325,	-0.625343, 0.000021	C	0.597701,	0.021997, -0.589343
N	-0.795784,	1.255096, 0.000000	N	0.182119,	-1.729258, 0.000007	N	0.618174,	1.314245, -0.293134
C	0.503943,	1.756599, 0.000000	C	1.523941,	-1.577347, -0.000026	C	-0.580271,	1.793147, 0.139750
C	1.573841,	0.939469, 0.000000	C	2.048139,	-0.306069, -0.000030	C	-1.687356,	0.991838, 0.295018
C	1.430213,	-0.517962, 0.000000	C	1.161895,	0.805799, 0.000002	C	-1.533115,	-0.389560, 0.026004
H	0.579870,	2.837735, 0.000000	H	2.137820,	-2.471990, -0.000049	H	-0.615975,	2.856525, 0.358897
H	2.577599,	1.345531, 0.000000	H	3.121159,	-0.154016, -0.000053	H	-2.631641,	1.401350, 0.637521
N	2.300773,	-1.383880, 0.000000	N	1.626245,	2.062699, 0.000018	N	-2.517065,	-1.284508, 0.227078
H	2.992163,	-2.099851, 0.000000	H	0.983365,	2.842923, 0.000074	H	-2.385435,	-2.227071, -0.119348
O	-2.258889,	-0.495552, 0.000000	H	2.617195,	2.262095, 0.000042	H	-3.461762,	-0.984103, 0.425661
H	-1.591207,	1.883222, 0.000000	O	-1.824712,	-1.050298, 0.000045	O	1.810705,	-0.507198, -1.032487
H	-0.179485,	-1.888689, 0.000000	H	-1.086205,	-2.167863, 0.000021	H	2.250353,	0.207948, -1.539339
Li	-3.842102,	-1.211785, 0.000000	Na	-2.677511,	1.040981, -0.000042	Na	2.486695,	-0.522007, 1.153536

Na ⁺ (C)-TS _{3,6}			Na ⁺ (C)-TS _{1,4}			Na ⁺ (C)-TS _{4,5}		
x, y, z			x, y, z			x, y, z		
N	0.666527,	-0.941756, 0.000072	N	0.354111,	-0.802479, 0.000008	N	0.666527,	-0.941756, 0.000072
C	-0.565839,	-0.381890, 0.000294	C	-0.546962,	0.201438, -0.000007	C	-0.565839,	-0.381890, 0.000294
N	-0.875868,	0.903473, 0.000310	N	-0.016821,	1.454639, 0.000000	N	-0.875868,	0.903473, 0.000310
C	0.229725,	1.707758, 0.000181	C	1.353470,	1.693517, 0.000004	C	0.229725,	1.707758, 0.000181
C	1.524739,	1.243791, 0.000005	C	2.245614,	0.685192, 0.000008	C	1.524739,	1.243791, 0.000005
C	1.759307,	-0.158429, -0.000101	C	1.817885,	-0.715663, 0.000003	C	1.759307,	-0.158429, -0.000101
H	0.034336,	2.776139, 0.000263	H	1.638815,	2.739283, 0.000006	H	0.034336,	2.776139, 0.000263
H	2.355969,	1.939452, -0.000032	H	3.309077,	0.888035, 0.000008	H	2.355969,	1.939452, -0.000032
N	2.973205,	-0.713476, -0.000614	N	2.506159,	-1.735174, -0.000008	N	2.973205,	-0.713476, -0.000614
H	3.081590,	-1.719988, 0.000438	H	3.062616,	-2.559431, 0.000017	H	3.081590,	-1.719988, 0.000438
H	3.809305,	-0.144721, 0.000566	O	-1.792286,	0.015954, -0.000025	H	3.809305,	-0.144721, 0.000566
O	-1.403442,	-1.394156, 0.000427	H	-0.674434,	2.224941, -0.000012	O	-1.403442,	-1.394156, 0.000427
Na	-3.174648,	0.096466, -0.000501	H	-0.014902,	-1.750009, -0.000011	Na	-3.174648,	0.096466, -0.000501
H	-0.167180,	-1.963829, 0.000208	Na	-3.827915,	-0.479660, 0.000013	H	-0.167180,	-1.963829, 0.000208

K ⁺ (C)TS _{1,2}			K ⁺ (C)-TS _{2,3}			K ⁺ (C)-TS _{3,6}		
x, y, z			x, y, z			x, y, z		
N	0.064443,	0.524155, 0.000174	N	0.737177,	-0.828333, 0.591401	N	-0.914455,	-0.004447, -1.112178
C	-0.039050,	-0.802828, 0.000089	C	-0.167225,	0.106201, 0.844053	C	-0.383273,	-0.000088, 0.122321
N	0.959664,	-1.706125, -0.000129	N	-0.224144,	1.367935, 0.428811	N	0.880299,	0.001965, 0.440935
C	2.232360,	-1.263935, -0.000132	C	0.877831,	1.771510, -0.257188	C	1.691019,	-0.000816, -0.642604
C	2.462074,	0.092018, -0.000024	C	1.928704,	0.929727, -0.541005	C	1.269943,	-0.005255, -1.939068
C	1.346851,	0.971905, 0.000053	C	1.813200,	-0.413215, -0.111601	C	-0.130630,	-0.007164, -2.183276
H	3.028412,	-2.001336, -0.000206	H	0.884311,	2.809651, -0.578481	H	2.744150,	0.000701, -0.431165
H	3.473718,	0.480562, -0.000073	H	2.797134,	1.280075, -1.088440	H	1.967904,	-0.007253, -2.752266
N	1.521635,	2.303494, -0.000027	N	2.733916,	-1.353270, -0.408328	N	-0.663633,	-0.011383, -3.393907
H	0.720394,	2.918877, 0.000182	H	2.666989,	-2.243357, 0.070564	H	-1.652030,	-0.012573, -3.515370
H	2.442912,	2.718109, 0.000085	H	3.640863,	-1.087685, -0.768030	H	-0.091750,	-0.013465, -4.207943
O	-1.147041,	-1.509509, 0.000235	O	-1.263281,	-0.338178, 1.558307	O	-1.400006,	0.001585, 0.900987
K	-2.850740,	0.449937, -0.000101	H	-1.596468,	0.439816, 2.050853	K	0.019192,	0.010506, 3.174813
H	-0.158656,	-2.422578, 0.000015	K	-2.512120,	-0.376986, -0.844298	H	-1.930715,	-0.002708, -0.310782

Table 6S. (continued) MP2(full)/6-31G* Optimized Geometries of Transition States for Unimolecular Tautomerization of Cytosine and M⁺(cytosine) Complexes.

K ⁺ (C)-TS _{1,4}			K ⁺ (C)-TS _{4,5}		
x, y, z			x, y, z		
N	0.639681,	0.714270, 0.000000	N	-0.799842,	-0.779466, 0.000000
C	-0.024477,	-0.451169, 0.000000	C	0.051206,	0.274551, 0.000000
N	0.799556,	-1.574538, 0.000000	N	-0.558754,	1.496637, 0.000000
C	2.169840,	-1.518089, 0.000000	C	-1.939164,	1.654203, 0.000000
C	2.836708,	-0.334461, 0.000000	C	-2.771728,	0.596169, 0.000000
C	2.018019,	0.836859, 0.000000	C	-2.260835,	-0.776491, 0.000000
H	0.323540,	-2.470414, 0.000000	H	-2.286280,	2.681282, 0.000000
H	2.678435,	-2.476424, 0.000000	H	-3.845218,	0.735533, 0.000000
H	3.918077,	-0.293285, 0.000000	N	-2.896272,	-1.831659, 0.000000
N	2.190798,	2.131302, 0.000000	H	-3.390776,	-2.693876, 0.000001
H	3.103502,	2.579418, 0.000000	H	1.298561,	0.159468, 0.000000
O	-1.266310,	-0.589652, -0.000001	O	0.051974,	2.303989, 0.000000
H	0.802483,	2.026843, 0.000000	H	-0.377022,	-1.703617, 0.000000
K	-3.584544,	0.276579, 0.000000	K	3.724529,	-0.278328, 0.000000

Table 7S. Enthalpies and Free Energies of Alkali Metal Ion Binding to Cytosine at 298 K in kJ/mol^a

System	ΔH_0	ΔH_0^b	$\Delta H_{298}-\Delta H_0^b$	ΔH_{298}	ΔH_{298}^b	$T\Delta S_{298}^b$	ΔG_{298}	ΔG_{298}^b
Li ⁺ (C ₁)	296.4 (6.6)	269.4	4.7 (0.3)	301.1 (6.6)	274.1	40.0 (0.6)	261.1 (6.6)	234.1
Li ⁺ (C ₂)	284.7 (6.7)	209.8	3.7 (0.4)	288.4 (6.7)	213.5	38.7 (0.5)	249.7 (6.7)	174.8
Li ⁺ (C ₃)	281.6 (6.9)	230.1	4.2 (0.4)	285.8 (6.9)	234.3	39.7 (0.5)	246.1 (6.9)	194.6
Li ⁺ (C ₄)	270.2 (7.3)	199.7	3.9 (0.3)	274.1 (7.3)	203.6	36.6 (0.5)	237.5 (7.3)	167.0
Li ⁺ (C ₅)		187.8	3.8 (0.3)		191.6	36.6 (0.5)		155.0
Li ⁺ (C ₆)		279.7	4.2 (0.5)		283.9	39.7 (0.5)		244.2
Na ⁺ (C ₁)	179.3 (4.5)	200.7	1.7 (0.2)	181.0 (4.5)	202.4	34.7 (0.6)	146.3 (4.5)	167.7
Na ⁺ (C ₂)	175.4 (4.8)	150.2	1.1 (0.3)	176.5 (4.8)	151.3	33.7 (0.7)	142.8 (4.8)	117.6
Na ⁺ (C ₃)	178.9 (6.2)	170.9	1.1 (0.3)	180.0 (6.2)	172.0	34.0 (0.7)	146.0 (6.2)	138.0
Na ⁺ (C ₄)	170.9 (4.8)	135.8	1.0 (0.1)	171.9 (4.8)	136.8	27.4 (0.5)	144.5 (4.8)	109.4
Na ⁺ (C ₅)		128.2	1.1 (0.2)		129.3	27.8 (0.5)		101.5
Na ⁺ (C ₆)		213.1	2.0 (0.2)		215.1	32.2 (0.6)		182.9
K ⁺ (C ₁)	138.2 (3.4)	158.9	1.9 (0.3)	140.1 (3.4)	160.8	37.7 (0.7)	102.4 (3.5)	123.1
K ⁺ (C ₂)	133.1 (3.5)	115.1	1.0 (0.1)	134.1 (3.5)	116.1	30.2 (0.5)	103.9 (3.5)	85.9
K ⁺ (C ₃)	136.0 (3.5)	133.7	0.3 (0.1)	133.4 (3.5)	134.0	32.5 (0.8)	101.5 (3.6)	101.5
K ⁺ (C ₄)	131.5 (3.8)	103.8	0.7 (0.2)	132.2 (3.8)	104.5	27.1 (0.6)	105.1 (3.8)	77.4
K ⁺ (C ₅)		96.5	0.6 (0.1)		97.1	27.2 (0.6)		69.9
K ⁺ (C ₆)		171.8	0.4 (0.2)		172.2	30.8 (0.7)		141.4

^aUncertainties are listed in the parentheses. ^bValues from calculations as described in text with frequencies scaled by 0.9646. Uncertainties in the enthalpic and entropic corrections are determined by 10% variation in the molecular constants.

Figure Captions

Fig. 1S. Cross sections for the collision-induced dissociation of $M^+(\text{cytosine})$ complexes, where $M^+ = \text{Li}^+$ and K^+ parts a and b respectively, with Xe as a function of collision energy in the center-of-mass frame (lower x-axis) and laboratory frame (upper x-axis). Data for the M^+ product channel are shown for a Xe pressure of 0.2 mTorr ($\bullet$) and extrapolated to zero ($\circ$).

Fig. 2S. Zero-pressure-extrapolated cross sections for collision-induced dissociation of $M^+(\text{cytosine})$ complexes, where $M^+ = \text{Li}^+$ and K^+ parts a and b respectively, with Xe in the threshold region as a function of kinetic energy in the center-of-mass frame (lower x-axis) and laboratory frame (upper x-axis). The solid lines show the best fits to the data using eq 1 convoluted over the neutral and ion kinetic and internal energy distributions. The dashed lines show the model cross sections in the absence of experimental kinetic energy broadening for reactants with an internal energy corresponding to 0 K.

Fig. 3S. RRKM rate constants for the unimolecular tautomerization, $M^+(\text{C}_x) \rightarrow M^+(\text{C}_y)$, where $(x,y) = (1,2), (3,6), (4,1), (3,2), (6,3), (1,4), (5,4),$ and $(4,5)$, parts a through h, respectively. The internal energy is provided by the association of M^+ with C_x , and taken from theoretical calculations at the MP2(full)/6-311+G(2d,2p)//MP2(full)/6-31G* level of theory including ZPE and BSSE corrections.

(41) Gaussian 98, Revision A.11, Frisch, M. J.; Trucks, G. W.; Schlegel, H. B.; Scuseria, G. E.; Robb, M. A.; Cheeseman, J. R.; Zakrzewski, V. G.; Montgomery Jr, J. A.; Stratmann, R. E.; Burant, J. C.; Dapprich, S.; Millam, J. M.; Daniels, A. D.; Kudin, K. N.; Strain, M. C.; Farkas, O.; Tomasi, J.; Barone, V.; Cossi, M.; Cammi, R.; Mennucci, B.; Pomelli, C.; Adamo, C.; Clifford, S.; Ochterski, J.; Petersson, G.A.; Ayala, P.Y.; Cui, Q.; Morokuma, K.; Salvador, P.; Dannenberg, J. J.; Malick, D.K.; Rabuck, A. D.; Raghavachari, K.; Foresman, J. B.; Cioslowski, J.; Ortiz, J. V.; Baboul, A. G.; Stefanov, B. B.; Liu, G.; Liashenko, A.; Piskorz, P.; Komaromi, I.; Gomperts, R.; Martin, R. L.; Fox, D. J.; Keith, T.; Al-Laham, M. A.; Peng, C. Y.; Nanayakkara, A.; Challacombe, M.; Gill, P. M. W.; Johnson, B.; Chen, W.; Wong, M. W.; Andres, J. L.; Gonzales, C.; Head-Gordon, M.; Replogle, E. S.; Pople, J. A.; Gaussian, Inc Pittsburgh, PA, 2001.

Figure 1S.

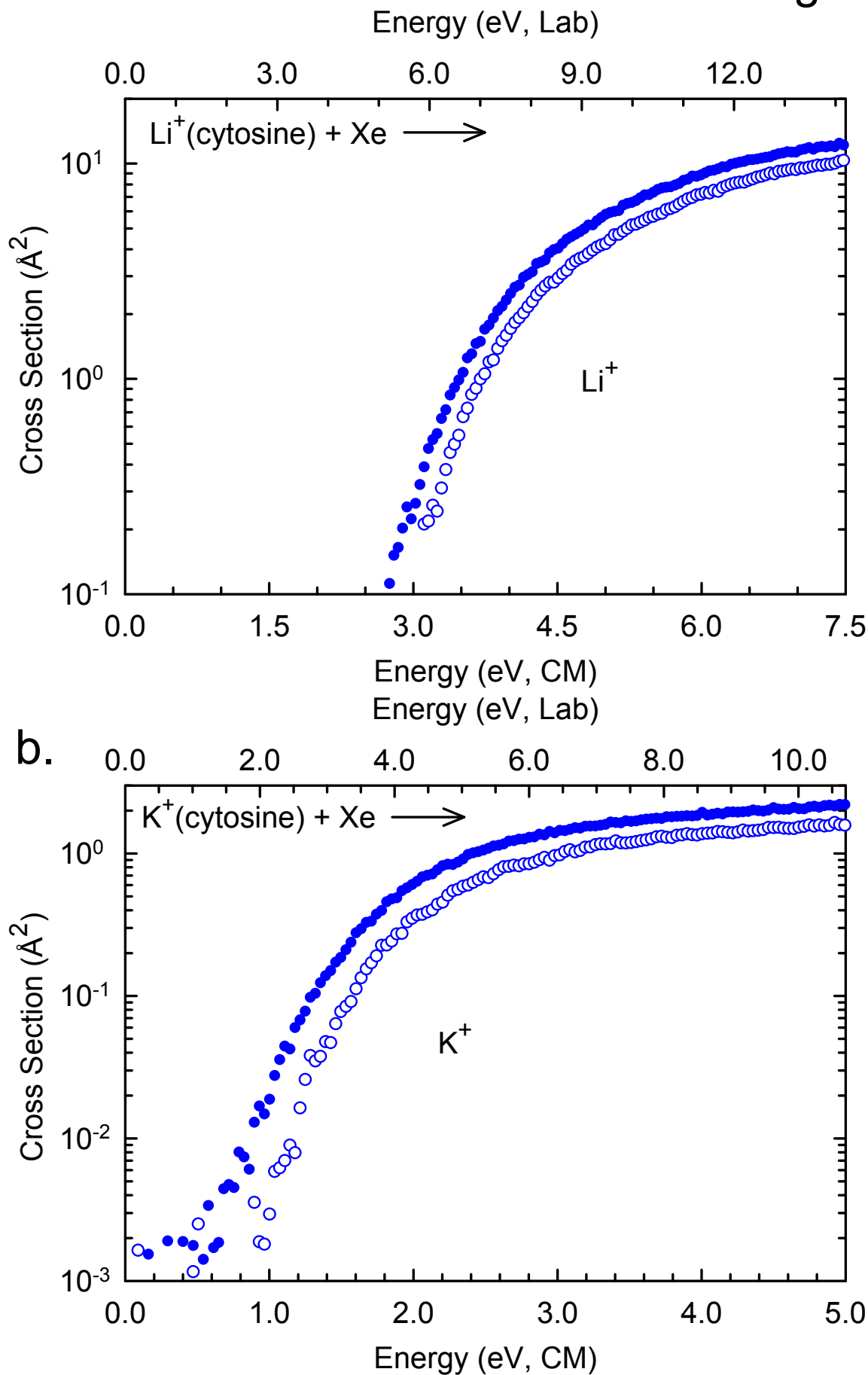


Figure 2S.

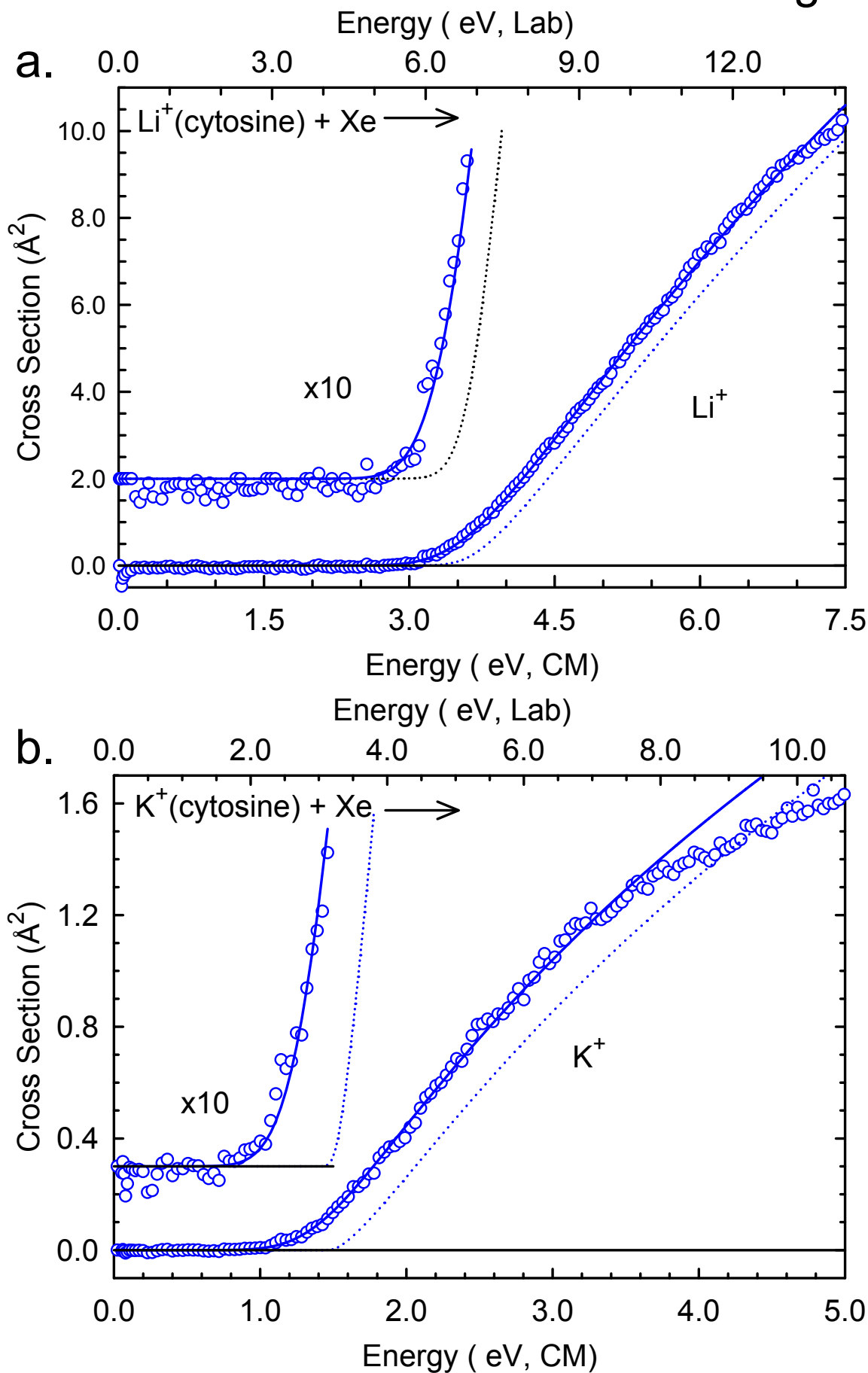


Figure 3S.

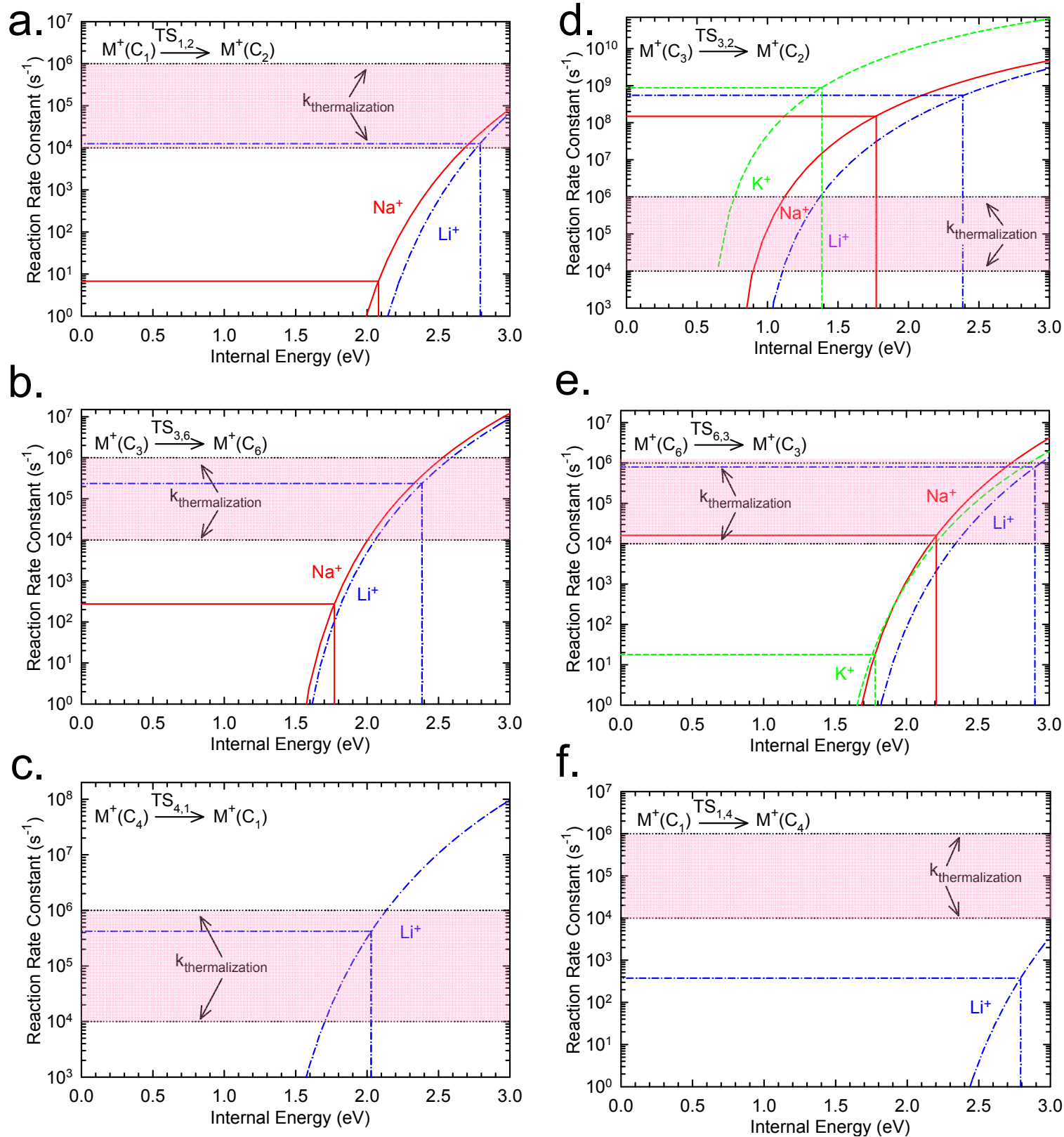


Figure 3S.

