Electronic Supplementary Information for ...

Structural Determination of Arginine-Linked Cisplatin Complexes via IRMPD Action Spectroscopy: Arginine Binds to Platinum via NO⁻-Binding Mode

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Figure Captions

Figure S1. IRMPD spectra of the $[(Arg-H)PtCl_2]^-$ and $[(Arg)PtCl_2+Na]^+$ complexes in the fingerprint region measured under variable attenuation of the FEL.

Figure S2. Binding modes of Arg to the Pt center of the [(Arg-H)PtCl₂]⁻ complex and the designated nomenclature.

Figure S3. Binding modes of Arg to the Pt center with possible sodium binding modes of the $[(Arg)PtCl_2+Na]^+$ complex and the designated nomenclature.

Figure S4. Stable low-energy conformers of the [(Arg–H)PtCl₂]⁻ complex and their relative Gibbs energies at 298 K calculated at B3LYP/mDZP/def2-TZVP level of theory. Structures are classified by the Arg binding mode to Pt.

Figure S5. Stable low-energy conformers of the $[(Arg)PtCl_2+Na]^+$ complex and their relative Gibbs energies at 298 K calculated at B3LYP/mDZP/def2-TZVP level of theory. Structures are classified by the Arg binding mode to Pt.

Figure S6. Comparison of the experimental IRMPD spectrum of $[(Arg-H)PtCl_2]^-$ with theoretical IR spectra predicted for select NN_s binding conformers along with their optimized structures and relative Gibbs energies computed at B3LYP/mDZP/def2-TZVP level of theory at 298 K. Misaligned IR features are highlighted in red.

Figure S7. Comparison of the experimental IRMPD spectrum of $[(Arg-H)PtCl_2]^-$ with theoretical IR spectra predicted for select NO⁻ binding conformers along with their optimized structures and relative Gibbs energies computed at B3LYP/mDZP/def2-TZVP level of theory at 298 K. Misaligned IR features are highlighted in red.

Figure S8. Comparison of the experimental IRMPD spectrum of $[(Arg-H)PtCl_2]^-$ with theoretical IR spectra predicted for select N_sO⁻ binding conformers along with their optimized structures and relative Gibbs energies computed at B3LYP/mDZP/def2-TZVP level of theory at 298 K. Misaligned IR features are highlighted in red.

Figure S9. Comparison of the experimental IRMPD spectrum of $[(Arg-H)PtCl_2]^-$ with theoretical IR spectra predicted for select OO⁻ binding conformers along with their optimized structures and relative Gibbs energies computed at B3LYP/mDZP/def2-TZVP level of theory at 298 K. Misaligned IR features are highlighted in red.

Figure S10. Comparison of the experimental IRMPD spectrum of $[(Arg-H)PtCl_2]^-$ with theoretical IR spectra predicted for select conformers with other side chain binding modes along with their optimized structures and relative Gibbs energies computed at B3LYP/mDZP/def2-TZVP level of theory at 298 K. Misaligned IR features are highlighted in red.

Figure S11. Comparison of the experimental IRMPD spectrum of $[(Arg)PtCl_2+Na]^+$ with theoretical IR spectra predicted for select NN_s binding conformers along with their optimized structures and relative Gibbs energies computed at B3LYP/mDZP/def2-TZVP level of theory at 298 K. Misaligned IR features are highlighted in red.

Figure S12. Comparison of the experimental IRMPD spectrum of $[(Arg)PtCl_2+Na]^+$ with theoretical IR spectra predicted for the NO⁻ binding conformers along with their optimized structures and relative Gibbs energies computed at B3LYP/mDZP/def2-TZVP level of theory at 298 K. Misaligned IR features are highlighted in red.

Figure S13. Comparison of the experimental IRMPD spectrum of $[(Arg)PtCl_2+Na]^+$ with theoretical IR spectra predicted for select N_sO binding conformers along with their optimized structures and relative Gibbs energies computed at B3LYP/mDZP/def2-TZVP level of theory at 298 K. Misaligned IR features are highlighted in red.

Figure S14. Comparison of the experimental IRMPD spectrum of $[(Arg)PtCl_2+Na]^+$ with theoretical IR spectra predicted for select OO⁻ binding conformers along with their optimized structures and relative Gibbs energies computed at B3LYP/mDZP/def2-TZVP level of theory at 298 K. Misaligned IR features are highlighted in red.

Figure S15. Comparison of the experimental IRMPD spectrum of $[(Arg)PtCl_2+Na]^+$ with theoretical IR spectra predicted for select conformers with other side chain binding modes along with their optimized structures and relative Gibbs energies computed at B3LYP/mDZP/def2-TZVP level of theory at 298 K. Misaligned IR features are highlighted in red.

Figure S1.



[(Arg-H)PtCl₂]⁻



 NN_s



NO







 NN_{ω}



 NN_{ϵ}





N_ωO⁻



N_εO⁻

[(Arg)PtCl₂+Na]⁺





NO









NO⁻_OCl_b





 $NN_sOCI_bCI_s$

 NN_sOCI_s







H₃



 NN_{ϵ}



 $NO_N_sCI_bCI_O$





 $N_{\epsilon}O$



ЪН H_2N

Ν_ωΟ



NH₂



NN_s_A 0.0 kJ/mol



 $NN_{s}E$ 14.9 kJ/mol



NN_s_I 31.2 kJ/mol



NN_s_M 38.3 kJ/mol



NN_s_Q 60.6 kJ/mol



NN_sB 7.4 kJ/mol



 $NN_{s}F$ 18.7 kJ/mol



NN_s_J 31.5 kJ/mol



NN_s_N



NN_s_R 66.1 kJ/mol



NN_s_C 12.6 kJ/mol



 $NN_{s}G$ 19.4 kJ/mol



 $NN_{s}K$ 33.2 kJ/mol



NN_s_O 41.5 kJ/mol







NN_s_P 43.0 kJ/mol

[(Arg-H)PtCl₂]⁻, NN_s



[(Arg-H)PtCl₂]⁻, NO⁻



[(Arg-H)PtCl₂]⁻, NO⁻

NO-_AR

24.2 kJ/mol

NO-_AV 25.1 kJ/mol



NO⁻_AO 22.7 kJ/mol



NO-_AS 24.3 kJ/mol



NO⁻_AW 25.3 kJ/mol



NO-_BA 26.0 kJ/mol



NO⁻_BE 26.3 kJ/mol



NO^{-_}AP 23.3 kJ/mol



NO⁻_AT 24.7 kJ/mol



NO⁻_AX 25.7 kJ/mol



NO⁻_BB 26.1 kJ/mol



NO⁻_BF 26.7 kJ/mol



23.9 kJ/mol



NO⁻_AU 24.9 kJ/mol



NO⁻_AY 25.9 kJ/mol



• NO⁻_BC 26.1 kJ/mol





NO⁻_BD 26.2 kJ/mol



[(Arg-H)PtCl₂]⁻, NO⁻



[(Arg-H)PtCl₂]⁻, NO⁻



A CAR

NO⁻_CC 36.4 kJ/mol







NO⁻_CI 40.0 kJ/mol

NO⁻_CJ 40.1 kJ/mol

NO-_CF

37.1 kJ/mol



NO-_CG

37.7 kJ/mol

NO⁻_CK 41.3 kJ/mol



NO-_CH

38.5 kJ/mol

NO⁻_CL 44.7 kJ/mol



NO⁻_CM 44.9 kJ/mol



NO⁻_CN 48.1 kJ/mol



S12





 $[(Arg-H)PtCl_2]^-$, other side chain binding modes



[(Arg)PtCl₂+Na]⁺, NN_s



NN_s_U 20.3 kJ/mol



NN_s_Y 22.6 kJ/mol

NN_s_AC 25.5 kJ/mol



NN_s_AG 27.1 kJ/mol



NN_s_AK 34.8 kJ/mol

NN_s_V 20.9 kJ/mol

NN_s_Z 22.9 kJ/mol

NN_s_AD 25.6 kJ/mol



NN_s_AH 29.3 kJ/mol



NN_s_AL

37.1 kJ/mol

NN_s_W 21.7 kJ/mol



NN_s_AA 23.5 kJ/mol



NN_s_AE 26.6 kJ/mol



NN_s_AI 34.1 kJ/mol



NN_s_AM 43.0 kJ/mol



NN_s_AJ 34.2 kJ/mol



NN_s_AN 46.5 kJ/mol









[(Arg)PtCl₂+Na]⁺, NN_s

Figure S5. **NO⁻_B** 2.8 kJ/mol **NO⁻_C** 7.7 kJ/mol **NO-_D** 7.8 kJ/mol **NO⁻_A** 1.8 kJ/mol **NO⁻_F** 11.1 kJ/mol **NO-_H** 12.1 kJ/mol NO-_E NO- G 9.7 kJ/mol 11.6 kJ/mol **NO⁻_I** 12.2 kJ/mol **NO⁻_L** 17.0 kJ/mol **NO⁻_K** 15.2 kJ/mol NO-_J 14.0 kJ/mol **NO⁻_N** 17.6 kJ/mol **NO-_O** 18.4 kJ/mol **NO⁻_M** 17.5 kJ/mol NO-_P 18.6 kJ/mol NO-_Q NO-_R NO-_S NO-_T 19.2 kJ/mol 20.0 kJ/mol 21.9 kJ/mol 24.2 kJ/mol

[(Arg)PtCl₂+Na]⁺, NO⁻

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[(Arg)PtCl₂+Na]⁺, NO⁻



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[(Arg)PtCl₂+Na]⁺, OO⁻



NO⁻_OCI_b_A 21.4 kJ/mol



NN_s_OCI_s_A 69.2 kJ/mol



N_sO⁻_OCI_s_A 97.7 kJ/mol



NN_s_OCI_bCI_s_A 36.7 kJ/mol



NN_ω_Α 80.1 kJ/mol



N_sO⁻_A 65.8 kJ/mol



NN_€_A 67.7 kJ/mol



NO_A 84.1 kJ/mol



NO_N_sCl_bCl_o_A 86.5 kJ/mol



121.2 kJ/mol



N_eO_A 145.9 kJ/mol

Figure S6.



Figure S7.



Figure S7.



Figure S8.



[(Arg-H)PtCl₂]⁻, N_sO⁻

Figure S9.



[(Arg-H)PtCl₂]⁻, OO⁻

Figure S10.



[(Arg–H)PtCl₂]⁻, Other side chain binding modes

Figure S11.



Figure S12.



[(Arg)PtCl₂+Na]⁺, NO⁻

Figure S13.



[(Arg)PtCl₂+Na]⁺, N_sO



[(Arg)PtCl₂+Na]⁺, OO⁻



[(Arg)PtCl₂+Na]⁺, Other side chain binding modes



[(Arg)PtCl₂+Na]⁺, Other side chain binding modes