

Computational Study of the Attack of $\cdot\text{OH}$ radicals on Aromatic Aminoacids

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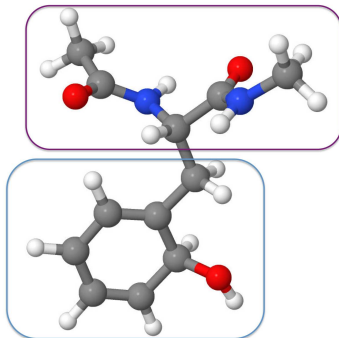
1. Introduction

ROS and biomacromolecules

- ROS: abundant among radical species (RNS, RCS, RSS)
 - $\cdot OH$, $\cdot OOH$, $\cdot O_2^-$, 1O_2 .
- Biomacromolecular targets:
 - Proteins, cell membrane phospholipids, DNA, RNA...
- Attack may cause...
 - Alteration of the function (function loss) of cell macromolecules
 - If not controlled, cell death
- Diseases
 - Neurodegenerative: Alzheimer, Parkinson etc...
 - Among others, diabetes mellitus, etc...

$\cdot\text{OH}$ Attack to Proteins

- Radical attack to the protein back-bone
 - Leads to breaking of back-bone
- Radical attack to the amino acid side chains
 - Leads to many different possibilities



Global Research Project

- Study reactivity with small models (DFT)
 - Attack to different positions of the back-bone:
 - Influence of the side chain important?
 - Influence of the protein folding?
(α -helix, β -sheet...)
 - Influence of the protein environment?
(different dielectrics)
 - Attack to different positions of the side chain
 - Influence of the protein folding?
 - Influence of the protein environment?
- Future QM/MM simulations on proteins

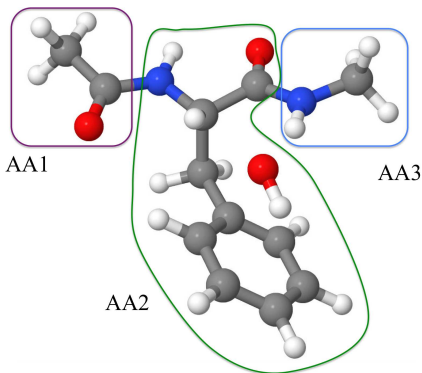
2. Model and Methods

Model

- Tripeptide model:
 - Two peptide bonds (α -helix, β -sheet...)
 - Side chain only at central amino acid.
 - Lateral amino acids truncated at C_α
- Disadvantages:
 - α -helix not properly described
- Advantages:
 - One side chain (R)

- This presentation

- R: Phe, Tyr, Trp
- β -sheet



Methods

- Geometry Optimizations & Frequency Calculations:
 - MPWB1K / 6-31+G(d,p)
- Singlepoints at $\epsilon = 4, 80$ (IEFPCM):
 - MPWB1K / 6-311++G(2df,p)
 - $H_{\epsilon}^{298} = E_{\epsilon} + H_{cont}^{gas}$
- Kinetics: $\Delta H_{\epsilon}^{TS} = H_{\epsilon}^{TS} - H_{\epsilon}^{React}$
- Thermodynamics: $\Delta H_{\epsilon}^{Prod} = H_{\epsilon}^{Prod} - H_{\epsilon}^{React}$
- Gaussian09

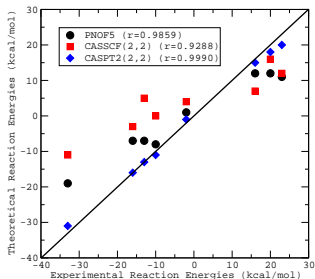
Method Validation

Reaction	Computed	Experimental	Difference
<i>Bond dissociation</i>			
$\text{CH}_4 \rightarrow \text{CH}_3^\cdot + \text{H}$	112.4	113.0	-0.6
$\text{NH}_3 \rightarrow \text{NH}_2^\cdot + \text{H}$	114.2	115.9	-1.7
$\text{H}_2\text{O} \rightarrow \cdot\text{OH} + \text{H}$	122.3	126.0	-3.7
$\text{C}_2\text{H}_6 \rightarrow \text{C}_2\text{H}_5^\cdot + \text{H}$	107.9	109.4	-1.5
$\text{H}_2\text{O}_2 \rightarrow \text{HOO}^\cdot + \text{H}$	89.1	92.7	-3.6
$\text{C}_2\text{H}_6 \rightarrow 2 \text{CH}_3^\cdot$	98.7	96.6	2.1
$\text{H}_2\text{O}_2 \rightarrow 2 \cdot\text{OH}$	49.4	55.0	-5.6
MAD			-2.1
<i>H abstraction</i>			
$\text{CH}_4 + \cdot\text{OH} \rightarrow \text{CH}_3^\cdot + \text{H}_2\text{O}$	-9.9	-13.0	3.1
$\text{NH}_3 + \cdot\text{OH} \rightarrow \text{NH}_2^\cdot + \text{H}_2\text{O}$	-8.2	-10.1	1.9
$\text{C}_2\text{H}_6 + \cdot\text{OH} \rightarrow \text{C}_2\text{H}_5^\cdot + \text{H}_2\text{O}$	-14.4	-16.6	2.2
$\text{H}_2\text{O}_2 + \cdot\text{OH} \rightarrow \text{HOO}^\cdot + \text{H}_2\text{O}$	-33.3	-33.3	0.0
MAD			1.8

Do not forget: DFT good results due to cancellation of errors!

PNOF5 and Radicals

- Energetics of radical mediated reactions
J. Chem. Theor. Comput. 8, 2646 (2012)

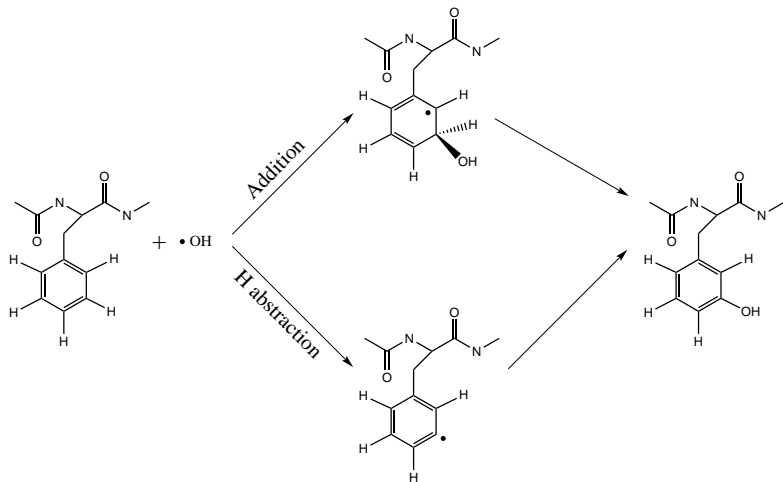


- Sign of reaction energetics improved from CASSCF to PNOF5

3. Results

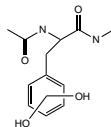
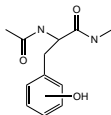
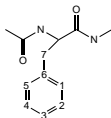
Mujika et. al., Chem. Eur. J 19, 6862 (2013)

Reactivity of $\cdot\text{OH}$ on Aromatic Rings

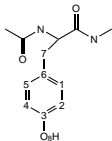


Atom Labeling and Experimental Products

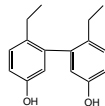
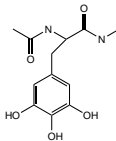
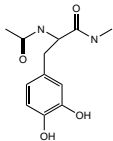
Phenylalanine



Tyrosine



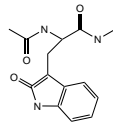
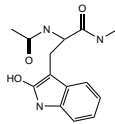
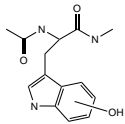
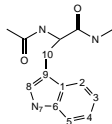
2-,3-, or 4-hydroxyphenylalanine



DOPA

Dityrosine

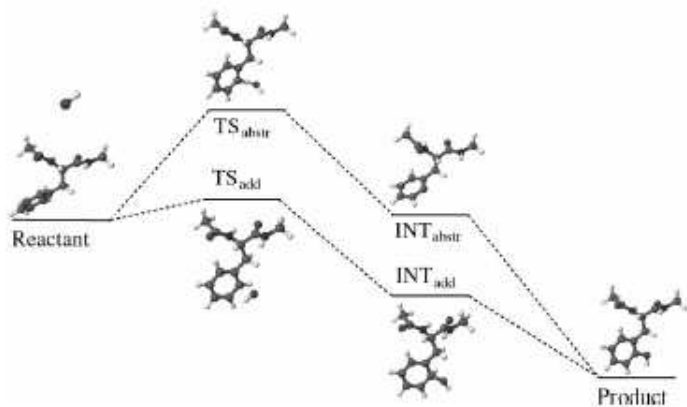
Tryptophan



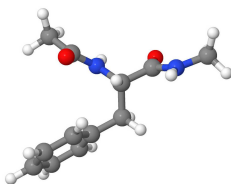
hydroxytryptophan

Oxindolylalanine

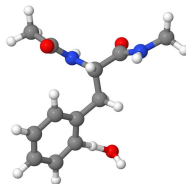
Phenylalanine



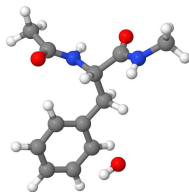
Phenylalanine



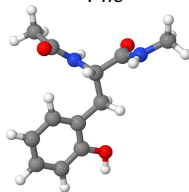
Reactant



TS^{C1ab}_{Phe}



TS^{C1add}_{Phe}



$Prod^{C1}_{Phe}$

Phenylalanine (1st ·OH)

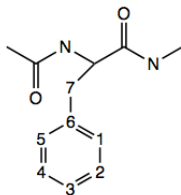
·OH Addition					H Abstraction				
	ΔH_4^{TS}	ΔH_4^{Int}	ΔH_{aq}^{TS}	ΔH_{aq}^{Int}		ΔH_4^{TS}	ΔH_4^{Int}	ΔH_{aq}^{TS}	ΔH_{aq}^{Int}
C1	-0.4	-18.0	1.1	-17.2	C1	4.1	-4.6	4.9	-4.9
C2	1.0	-16.3	1.7	-16.0	C2	3.7	-4.7	4.1	-5.5
C3	0.4	-18.0	1.0	-17.0	C3	4.1	-4.2	4.7	-4.9
C4	1.0	-16.5	1.7	-15.5	C4	5.3	-5.4	6.7	-6.2
C5	-4.5	-17.7	-1.3	-15.1	C5	2.2	-4.9	5.4	-5.4
C6	-0.5	-18.6	0.6	-14.9	C7	0.1	-29.7	3.2	-29.1

- $\epsilon=4$, 80 similar conclusions.
- Addition kinetically and thermodynamically favored
- Exception: H abstraction from C _{β}

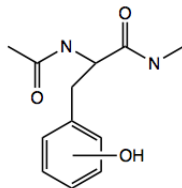
Phenylalanine (2nd ·OH)

	ΔH_4^{298}
Prod _{Phe} ^{C1}	-117.1
Prod _{Phe} ^{C2}	-116.9
Prod _{Phe} ^{C3}	-116.5
Prod _{Phe} ^{C4}	-116.4
Prod _{Phe} ^{C5}	-115.9
Prod _{Phe} ^{C7}	-111.0
Prod _{Phe} ^{C7α}	-102.5

Labeling



Exper. Products



2-,3-, or 4-hydroxyphenylalanine

- Addition + Abstraction
- Steric effects

Tyrosine ($1^{\text{st}} \cdot\text{OH}$)

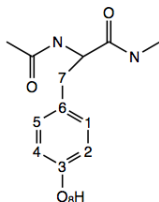
$\cdot\text{OH}$ Addition			H Abstraction		
	ΔH_4^{TS}	ΔH_4^{Int}		ΔH_4^{TS}	ΔH_4^{Int}
C1	0.9	-16.7	C1	5.2	-3.7
C2	0.6	-18.1	C2	6.2	-2.2
C3	2.5	-18.9	C4	5.8	-2.6
C4	1.1	-18.2	C5	3.0	-3.8
C5	-3.4	-17.0	C7	0.9	-29.6
C6	-1.8	-19.5	O8	2.4	-29.3

- Addition kinetically and thermodynamically favored
- Exception: H abstraction from C_β and O8

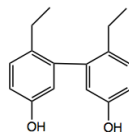
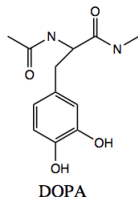
Tyrosine (2nd ·OH)

	ΔH_4^{298}
Prod _{Tyr} ^{C1}	-116.5
Prod _{Tyr} ^{C2}	-114.9
Prod _{Tyr} ^{C4}	-114.1
Prod _{Tyr} ^{C5}	-116.4
Prod _{Tyr} ^{C7}	-111.0
Prod _{Tyr} ^{C7α}	-101.3
Prod _{Tyr-Tyr} ^{OO}	-37.3
Prod _{Tyr-Tyr} ^{CC}	-107.7

Labeling



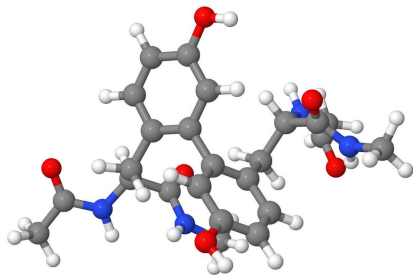
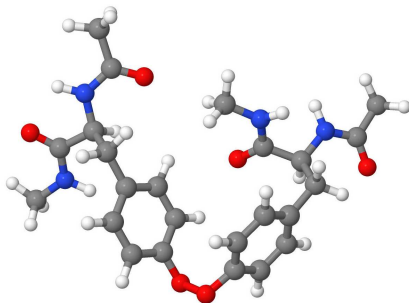
Exper. Products



Dityrosine

- DOPA: Addition + Abstraction
- Dityrosine: 2 H abstractions
- Thermodynamic effects

Tyrosine ($2^{nd} \cdot OH$)



Tryptophan ($1^{\text{st}} \cdot\text{OH}$)

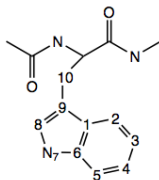
$\cdot\text{OH}$ Addition			H Abstraction		
	ΔH_4^{TS}	ΔH_4^{Int}		ΔH_4^{TS}	ΔH_4^{Int}
C2	0.1	-24.4	C2	1.7	-4.4
C3	0.7	-15.3	C3	4.7	-3.1
C4	-0.4	-17.7	C4	4.9	-3.3
C5	-0.8	-19.8	C5	5.8	-2.1
C8	-5.7	-28.6	N7	2.2	-24.5
			C8	-	3.3
			C10	-1.6	-28.7

- Addition kinetically and thermodynamically favored
- Exception: H abstraction from C_β and N7

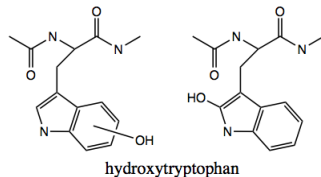
Tryptophan ($2^{nd} \cdot OH$)

	ΔH_4^{298}
Prod $_{Trp}^{C2}$	-116.2
Prod $_{Trp}^{C3}$	-113.9
Prod $_{Trp}^{C4}$	-114.6
Prod $_{Trp}^{C5}$	-114.6
Prod $_{Trp}^{N7}$	-74.7
Prod $_{Trp}^{C8}$	-123.5
Prod $_{Trp}^{C10}$	-111.6
Prod $_{Trp}^{10\alpha}$	-104.6

Labeling

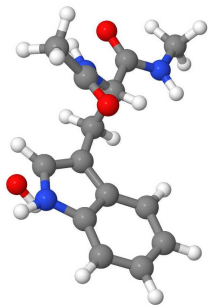


Exper. Products

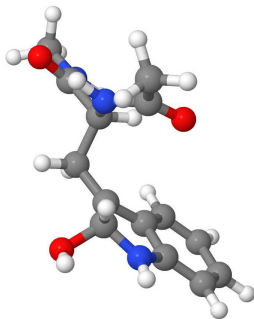


- Addition + Abstraction
- Thermodynamic effects

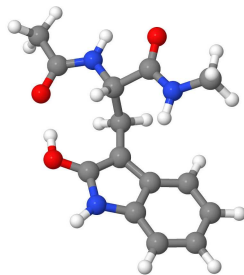
Tryptophan



TS_{Trp}^{C8add}



Int_{Trp}^{C8add}



$Prod_{Trp}^{C8}$

4. Conclusions

Conclusions

- All aromatic amino acids target.
- Thermodynamics governs over the kinetics
- Other factors, radical concentration and steric effects, important
- Two-step mechanism:
 - the first $\cdot\text{OH}$ radical addition
 - the second $\cdot\text{OH}$ radical abstracts a hydrogen.
- Results in agreement with experiments

Acknowledgments

Theoretical Chemistry Group



- The SGI/IZO-SGIker (UPV/EHU)
 - Arina supercomputer
- Funding
 - Basque and Spanish Governments
- Organizing committee
- Audience!