### **Supporting Information**

# Highly Efficient Ru(II)-Alkylidene Based Hoveyda-Grubbs Catalysts for Ring-Closing Metathesis Reactions

Mariam Y. Al-Enezi, Elizabeth John, Yehia A. Ibrahim and Nouria A. Al-Awadi\*

\*Chemistry Department, Kuwait University, P.O. Box 5969, Safat 13060, Kuwait. Email: <u>na.nabaq.kw@gmail.com</u>

#### **Table of Contents**

NMR Spectra	2-22
Mass spectra	22-30
Deconvoluted HRMS (ESI)	30-39
Single Crystal X-ray Diffraction Studies of the Ru complexes <b>7a</b> , <b>7b</b> & <b>7c</b>	40-42

#### NMR spectra



Figure S2. <sup>13</sup>C-NMR spectrum of **6a** in DMSO-d<sub>6</sub> at 25  $^{\circ}$ C.



Figure S3. <sup>1</sup>H-NMR spectrum of **6b** in CDCl<sub>3</sub> at 25  $^{\circ}$ C.



Figure S4. <sup>13</sup>C-NMR spectrum of **6b** in CDCl<sub>3</sub> at 25  $^{\circ}$ C.



Figure S6.  $^{13}$ C-NMR spectrum of 6c in CDCl<sub>3</sub> at 25  $^{0}$ C.



























Figure S15. <sup>1</sup>H-NMR spectrum of 8k in CDCl<sub>3</sub> at 25 <sup>0</sup>C.







Figure S17. 19F NMR spectrum of 8k in CDCl<sub>3</sub> at 25 <sup>o</sup>C.













Figure S23. 19F NMR spectrum of 8q in CDCl<sub>3</sub> at 25 <sup>o</sup>C.















Figure S28. <sup>1</sup>H-NMR spectrum of 9j in CDCl<sub>3</sub> at 25 <sup>0</sup>C.















Figure S34. <sup>1</sup>H-NMR spectrum of **90** in CDCl<sub>3</sub> at 25  $^{0}$ C.



Figure S36. 19F NMR spectrum of 9o in CDCl<sub>3</sub> at 25 <sup>o</sup>C.











Figure S39. 19F NMR spectrum of 9q in CDCl<sub>3</sub> at 25 <sup>o</sup>C.





















Figure S46. MALDI-MS spectrum of 7b.

BG FAB-782-re #16-20 RT: 0.67-0.84 AV: 5 NL: 2.49E6 T: + c FAB Full ms [49.50-900.50]







Figure S48. El spectrum of 8i.







Figure S50. El spectrum of 8o.







Figure S54. El spectrum of 9j.



Figure S56. El spectrum of 9m.









#### **Deconvoluted HRMS (ESI)**









Figure S62. HRMS spectrum of 6c.







Figure S66. HRMS spectrum of 8k.











Figure S72. HRMS spectrum of 9I.





HRMS-DFG-cmass1 #119 RT: 10.35 AV: 1 NL: 5.25E3 T: + c EI Full ms [149.50-220.50]













#### Single Crystal X-ray diffraction study of the Ru complexes 7a, 7b & 7c

Single crystals of **7a**, **7b & 7c** were grown by solvent diffusion method. The single crystal data collection of **7a** and **7c** were made on Bruker X8 prospector diffractometer by Cu-Kα. The reflection frames were then integrated with the Bruker SAINT Software package using a narrow-frame algorithm; the structures were solved using the Bruker SHELXTL Software Package. The single crystal data collection of **7b** was made on Rigaku Rapid II diffractometer by Mo-Kα This structure was then solved by 'crystalstructure' software package. Finally, all structures were refined using SHELXL-2017/1. All non-hydrogen atoms were refined anisotropically and hydrogen atoms were refined using the riding model. The crystal structures (thermal ellipsoid representation) of **7a**, **7b & 7c** are given in **Figure S72-S74** and their various crystallographic data including crystal nature, data collection strategy and refinement parameters are summarized in the **Table S1**.



Figure S72. crystal structure (thermal ellipsoid representation; 50% probability) of 7a.



Figure S73. crystal structure (thermal ellipsoid representation; 50% probability) of 7b.



Figure S74. crystal structure (thermal ellipsoid representation; 50% probability) of 7c.

Crystal sample	7a	7b	7c
Chemical formula	$C_{38}H_{43}CI_2N_3O_3Ru$	$C_{44}H_{49}CI_4N_3O_4RuS$	$C_{37}H_{43}Cl_2N_3O_3RuS$
M <sub>r</sub>	761.72	958.79	781.77
Crystal system, space group	Triclinic, P-1	Monoclinic, P2 <sub>1</sub> /c	Trigonal, R -3 :H
Temperature (K)	296	150	150
<i>a, b, c</i> (Å)	8.8611 (4), 10.6837 (5),	12.038 (10), 14.886 (12),	28.6420 (5), 24.4463 (9)
	19.3949 (10)	24.180 (19)	
α, β, γ (°)	74.511 (2), 87.898 (2), 89.558 (2)	90, 91.36 (3), 90	90, 90, 120
V (Å <sup>3</sup> )	1768.22 (15)	4332 (6)	17368.0 (9)
Z	2	4	18
Radiation type	Cu <i>Κ</i> α	Μο Κα	Cu <i>Κ</i> α
μ (mm <sup>-1</sup> )	5.30	0.70	5.36
Crystal size (mm)	0.21 × 0.20 × 0.08	0.10 × 0.05 × 0.04	0.18 × 0.14 × 0.12
Diffractometer	Bruker APEX-II CCD	Rigaku R-AXIS RAPID	Bruker APEX-II CCD
Absorption correction	Multi-scan	Multi-scan	Multi-scan
	SADABS2016/2 - Bruker AXS area detector scaling and absorption correction	ABSCOR (Rigaku, 1995)	SADABS2016/2 - Bruker AXS area detector scaling and absorption correction
T <sub>min</sub> , T <sub>max</sub>	0.220, 0.495	0.379, 0.972	0.45, 0.57
No. of measured, independent & observed $[l > 2\sigma(l)]$ reflections	37958, 6204, 5934	24863, 7781, 3493	61572, 6025, 4870
R <sub>int</sub>	0.040	0.171	0.067
(sin θ/λ) <sub>max</sub> (Å <sup>-1</sup> )	0.595	0.602	0.598
<i>R</i> [F <sup>2</sup> > 2σ(F <sup>2</sup> )], wR(F <sup>2</sup> ), S	0.027, 0.069, 1.09	0.077, 0.214, 0.96	0.053, 0.151, 1.03
No. of reflections	6204	7781	6025
No. of parameters	431	515	431
H-atom treatment	Constrained	Constrained	Constrained
Δρ <sub>max</sub> , Δρ <sub>min</sub> (e Å <sup>-3</sup> )	0.40, -0.29	1.01, -0.75	0.74, -0.51

## Table S1. Summary on the nature and various crystallographic parameters of 7a, 7b & 7c.