Electronic Supplementary Information

Metal Center and Aromatic Moiety in Schiff Base Complexes: Impact on G-Quadruplex Stabilization and Oncogene Downregulation.

Aurane Froux,^{a,b} Luisa D'Anna,^a Aurianne Rainot,^{a,c} Camille Neybecker,^b Angelo Spinello,^a Riccardo Bonsignore,^a Raphaël Rouget,^b Guillaume Harlé,^b Alessio Terenzi,^a Antonio Monari,^c Stéphanie Grandemange,^{b,*} and Giampaolo Barone,^{a,*}

a Università di Palermo, Department of Biological, Chemical and Pharmaceutical Sciences and Technologies, Viale delle Scienze, Edificio 17, 90128 Palermo, Italy

b Université de Lorraine and CNRS, UMR 7039 CRAN, F-54000 Nancy, France

c Université Paris Cité and CNRS, ITODYS, F-75006 Paris, France



Figure S1. Structures of the complexes synthetized in this study.



Figure S2. ¹H-NMR spectra of the new pyridinium (B) and metal complexes 1, 2, 4, 5, 6, 7, 8, 10 and 11 in d_6 -DMSO.













Figure S3. Mass spectra of compound **B** and metal complexes **1-12.** In each figure, calculated values (table), theorical spectrum (left) and experimental spectrum (right) are reported.

Compound ${\bm B}$ Calculated for $C_{14}H_{14}NO_2$ [2M + Cl]+ 491.17; found 491.17



Compound ${\rm 1\!\!I}$ Calculated for $(C_{34}H_{30}N_4O_2Ni)_2(ClO_4)_2\,[M]^{2+}\,684.12;$ found 684.12



Compound **1** Calculated for C₂₂H₁₆N₂NiO₂ [M]²⁺ 199.03; found 199.03



m/z	Abund (% largest)	Abund (% sum)	(% first)
199,03	100	52,98	100
199,53	24,79	13,13	24,79
200,03	41,87	22,19	41,87
200,53	11,55	6,12	11,55
201,03	7,07	3,75	7,07
201,53	1,51	0,8	1,51
202,02	1,55	0,82	1,55
202,53	0,36	0,19	0,36
203,03	0,05	0,02	0,05
203,53	0	0	0





94,07 100 93,32 100 95,07 6,95 6,48 6,95 96,07 0,21 0,19 0,21 97,07 0 0 0	93,32 100 6,48 6,95 0,19 0,21 0 0	07 100 07 6.95 07 0.21 07 0
95.07 6.95 6.48 6.95 96.07 0.21 0.19 0.21 97.07 0 0 0	6,48 6,95 0,19 0,21 0 0	07 6.95 07 0.21 07 0
96,07 0,21 0,19 0,21 97,07 0 0 0	0.19 0.21 0 0	07 0.21 07 0
97,07 0 0 0	0 0	07 0



Calculated

Compound **2** Calculated for $(C_{34}H_{30}N_4O_2Zn)_2(ClO_4)_2$ [M]²⁺ 691.11; found 691.11



Compound ${\bf 2}$ Calculated for $C_{22}H_{16}N_2ZnO_2\,[M]^{2+}$ 202.02; found 202.02



m/z	(% largest)	(% sum)	(% first)
202,02	100	37,57	100
202,53	24,79	9,31	24,79
203,02	61,31	23,03	61,31
203,52	23,19	8,71	23,19
204,02	43,49	16,34	43,49
204,52	10,24	3,85	10,24
205,02	2,67	1	2,67
205,53	0,45	0,17	0,45
206,03	0,05	0.02	0.05





m/z	Abund (% largest)	Abund (% sum)	Abund (% first)
94,07	100	93,32	100
95,07	6,95	6,48	6,95
96,07	0,21	0,19	0.21
07.07	0	0	0



Calculated

Compound **3** Calculated for $(C_{34}H_{30}N_4O_2Cu)_2(ClO_4)_2$ [M]²⁺ 689.11; found 689.12



Compound **3** Calculated for $C_{22}H_{16}N_2CuO_2$ [M]²⁺ 201.52; found 201.53



m/z	(% largest)	(% sum)	(% first)
201,52	100	53,82	100
202,03	24,79	13,34	24,79
202,52	47,97	25,81	47.97
203,03	11,38	6,13	11,38
203,53	1.52	0.82	1.52
204,03	0,15	0.08	0.15
204,53	0.01	0,01	0.01
205,03	0	0	0
205 53	0	0	0





Compound **4** Calculated for $(C_{34}H_{30}N_4O_2Pd)_2(ClO_4)_2$ [M]²⁺ 732.09; found 732.09



Compound ${\bf 4}$ Calculated for $C_{22}H_{16}N_2PdO_2\,[M]^{2+}$ 223.01; found 223.01



m/z	Abund (% largest)	Abund (% sum)	Abund (% first)
221,01	3,07	0,79	100
221,51	0,76	0,2	24,79
222,01	33,62	8,7	1095,51
222,51	75,5	19,53	2460,24
223,01	100	25,87	3258,66
223,51	22,74	5,88	741.04
224,01	82,59	21,36	2691,31
224,51	20.01	5,18	652,2
225,01	37,95	9,82	1236,68
225,51	9	2,33	293,24
226,02	1,2	0,31	39,16
226,52	0,12	0.03	3,76
227,02	0.01	0	0,28



Calculated for C_6H_7N [M + H]⁺ 94.07; found 94.07



m/z	Abund (% largest)	Abund (% sum)	Abund (% first)
94.07	100	93,32	100
95,07	6,95	6,48	6,95
96,07	0,21	0,19	0,21
97 07	0	0	0



Calculated

Found

Compound **5** Calculated for $(C_{34}H_{30}N_4O_2Pt)_2(Cl)_2 \ [M]^{2+}$ 757.17; found 757.17



Compound ${\bf 5}$ Calculated for $C_{22}H_{16}N_2PtO_2\,[M]^{2+}$ 267.54; found 267.54



m/z	Abund (% largest)	Abund (% sum)	Abund (% first)
265,04	0.03	0.01	100
265,54	0,01	0	24,79
266,04	1,86	0,61	5589,07
266,54	0.46	0,15	1384,77
267.04	78,54	25,68	235665,88
267,54	100	32,69	300039,72
268,04	82,69	27,03	248091,8
268,54	17,85	5,84	53553,36
269,04	19,35	6,33	58050,08
269,55	4,44	1,45	13328,6
270.05	0.59	0,19	1763,34
270,55	0,06	0.02	168,78
271.05	0	0	12.69



Calculated for $C_6H_7N [M + H]^+94.07$; found 94.07







Calculated

Compound **6** Calculated for $(C_{30}H_{30}N_4O_2Ni)_2(ClO_4)_2\,[M]^{2+}\,636.12$; found 636.12



Compound **6** Calculated for $C_{18}H_{16}N_2NiO_2$ [M]²⁺ 175.03; found 175.03

Ŧ					
ŧ					
Ť					
1					
0.8					
Ŧ					
0.7 +					
ŧ					
0.6 +					
+					
0.6					
Ŧ					
0.4	1				
ŧ					
0.3 I					
ŧ					
Ī					
÷					
0.1 +		1.			
Ŧ					

m/z	(% largest)	(% sum)	(% first)
175,03	100	55,31	100
175,53	20,46	11,32	20,46
176,03	40,92	22,63	40.92
176,53	9,76	5,4	9,76
177,02	6,62	3,66	6,62
177,53	1,21	0,67	1,21
178,02	1,5	0,83	1.5
178,53	0,29	0,16	0,29
179,03	0.03	0.02	0.03





94.07 100 93.32 100 35.07 6.95 6.48 6.95 36.07 0.21 0.19 0.21 37.07 0 0 0	94,07 100 93,32 100 95,07 6,95 6,48 6,95 96,07 0,21 0,19 0,21 97,07 0 0 0 0	94,07 100 93,32 100 95,07 6,95 6,48 6,95 96,07 0,21 0,19 0,21 97,07 0 0 0	94.07 100 93.32 100 95.07 6.95 6.48 6.95 96.07 0.21 0.19 0.21 97.07 0 0 0	m/z	Abund (% largest)	Abund (% sum)	Abund (% first)
35.07 6.95 6.48 6.95 36.07 0.21 0.19 0.21 37.07 0 0 0	95.07 6.95 6.48 6.95 96.07 0.21 0.19 0.21 97.07 0 0 0 0	95.07 6.95 6.48 6.95 96.07 0.21 0.19 0.21 97.07 0 0 0 0	95.07 6,95 6,48 6,95 96.07 0.21 0.19 0.21 97.07 0 0 0	94,07	100	93,32	100
36.07 0.21 0.19 0.21 37.07 0 0 0	96.07 0.21 0.19 0.21 97.07 0 0 0 0	96.07 0.21 0.19 0.21 97.07 0 0 0 0	96.07 0.21 0.19 0.21 97.07 0 0 0 0	95,07	6,95	6,48	6,95
97,07 0 0 0	97.07 0 0 0	97,07 0 0 0	97,07 0 0 0	96,07	0,21	0,19	0,21
				97 07	0	0	0
					U	U	0



Compound 7 Calculated for $C_{18}H_{16}N_2ZnO_2 [M]^{2+}$ 178.02; found 178.02



m/z	(% largest)	Abund (% sum)	Abund (% first)
178,02	100	39,22	100
178,53	20,46	8,02	20,46
179,02	60,35	23,67	60,35
179,52	20,56	8,06	20,56
180,02	42,56	16,69	42,56
180,52	8,39	3,29	8,39
181,02	2,28	0,89	2,28
181,52	0,35	0,14	0.35
182,03	0.04	0,01	0.04
182,53	0	0	0





Compound **8** Calculated for $C_{18}H_{16}N_2CuO_2$ [M]²⁺ 177.52; found 177.53



m/z	Abund (% largest)	Abund (% sum)	Abund (% first)
177,52	100	56,18	100
178,03	20,46	11,49	20,46
178,52	47,01	26,41	47.01
179,03	9,33	5,24	9,33
179,53	1,08	0,61	1.08
180,03	0.09	0.05	0.09
180 53	0.01	0	0.01







Compound $\,{\rm 9}$ Calculated for $C_{18}H_{16}N_2PdO_2\,[M]^{2+}$ 199.01; found 199.01



m/z	Abund (% largest)	Abund (% sum)	Abund (% first)
197.01	3,17	0,83	100
197,51	0.65	0,17	20.46
198,01	34,71	9,07	1094,55
198,51	76,51	20	2412,87
199,01	100	26,13	3153,5
199,51	19,12	5	602,91
200,01	84.45	22,07	2663
200,51	17.01	4,45	536,56
201.01	38,42	10,04	1211,59
201.51	7.62	1,99	240,43
202,02	0,88	0,23	27.9
202,52	0.08	0.02	2.38
203,02	0.01	0	0.16
203,52	0	0	0.01



Calculated for $C_6H_7N [M + H]^+94.07$; found 94.08



Compound $\, {\rm 10}\,$ Calculated for $C_{26}H_{18}N_2NiO_2\, [M]^{2+}$ 224.04; found 224.03



m/z	Abund (% largest)	Abund (% sum)	Abund (% first)	
224,04	100	50,74	100	
224,54	29,13	14,78	29,13	
225,03	43,02	21,83	43,02	
225,53	13,39	6,79	13,39	
226,03	7,6	3,86	7,6	
226,53	1,82	0,92	1,82	
227,03	1,62	0,82	1,62	
227,53	0,42	0,22	0,42	
228,04	0,06	0,03	0,06	
228,54	0.01	0	0,01	









Compound ~11 Calculated for $C_{26}H_{18}N_2ZnO_2~[M]^{2+}~227.03;$ found 227.03



m/z	Abund (% largest)	Abund (% sum)	Abund (% first)	3
227,03	100	35,98	100	
227,53	29,13	10,48	29,13	
228,03	62,46	22,47	62,46	
228,53	25,87	9,31	25,87	_
229,03	44,54	16,02	44,54	_
229,53	12,15	4,37	12,15	
230,03	3,15	1,13	3,15	
230,53	0,58	0,21	0.58	
231,03	0.08	0.03	0.08	-
231,54	0.01	0	0.01	



Calculated for C_6H_7N [M + H]⁺94.07; found 94.08



m/z	Abund (% largest)	Abund (% sum)	Abund (% first)
94,07	100	93,32	100
95,07	6,95	6,48	6,95
96,07	0,21	0,19	0,21
97,07	0	0	0



Compound **12** Calculated for $(C_{38}H_{32}N_4O_2Cu)_2(ClO_4)_2$ [M]²⁺ 739.13; found 739.13



Compound ~12 Calculated for $C_{26}H_{18}N_2CuO_2\,[M]^{2\star}$ 226.53; found 226.53



m/z	Abund (% largest)	Abund (% sum)	Abund (% first)
226,53	100	51,54	100
27,03	29,13	15,02	29,13
27,53	49,11	25,31	49,11
28,03	13,49	6,95	13,49
28,53	2,05	1,06	2.05
29,04	0,22	0,11	0,22
29,54	0,02	0,01	0.02
30,04	0	0	0
30,54	0	0	0





x10	Abund (% first)	Abund (% sum)	Abund (% largest)	m/z
	100	93,32	100	94,07
7	6,95	6.48	6,95	95,07
6	0.21	0.19	0.21	96,07
	0	0	0	97,07
5				
4				
4				
4				
4				
4 3 2				
4 3 2				
4 3 2 1				
4 3 2 1				



Figure S4. Example of FRET melting profile of cMYC G4 (0.2 μ M, black line) upon interaction with compound **1** (1 μ M, red line) in a [G4 DNA]:[compound **1**] ratio of 1:5 in 10 mM potassium cacodylate buffer, pH 7.2, showing that in such conditions the melting temperature cannot be reached.







Figure S6. UV-vis spectra of compound (1) at 20 μ M in presence of increasing concentration of ctDNA, *kRAS*, *BCL2* or *cMYC* quadruplexes in 50 mM Tris HCl 100 mM KCl pH 7.4.







Figure S8. UV-vis spectra of compound (10) at 20 μ M in presence of increasing concentration of ctDNA, *kRAS*, *BCL2* or *cMYC* quadruplexes in 50 mM Tris HCl 100 mM KCl pH 7.4.



Figure S9. IC50 determination of metal complexes 1 – **12 on T3M4 cells.** Relative viability (%) of T3M4 cell line at 48 hours of treatment of the considered complex metal, obtained by crystal violet assay. *Mean* \pm *SEM* (n=5).



Figure S10. IC50 determination of metal complexes 1 – 12 on T47D cells. Relative viability (%) of T47D cell line at 48 hours of treatment of the considered complex metal, obtained by crystal violet assay. *Mean* \pm *SEM* (n=5).



Figure S11. IC50 determination of metal complexes 1 - 12 on MDA-MB-231 cells. Relative viability (%) of MDA-MB-231 cell line at 48 hours of treatment of the considered complex metal, obtained by crystal violet assay. *Mean* ± *SEM* (n=5).



Figure S12. Doubling time of T3M4, T47D and MDA-MB-231 cell lines. Doubling time of T3M4 (A), T47D (B), and MDA-MB-231 (C) cell lines was determined from the proliferation follow-up of $t \ln 2$

untreated cells over 72 hours using the equation: $T = \overline{\ln(day 1) - \ln(day 2)}$. Mean $\pm SEM$ (n=6).



Table S1. DNA sequences (5' - 3') of the G4 motifs used for the DNA binding studies. The oligonucleotides were used free for the UV-visible and Circular Dichroism assays, and fluorolabelled with FAM and TAMRA probes for FRET experiments. dsDNA was used only for the FRET experiment.

Oligonucleotide	Sequence
dsDNA	5'-/56-FAM/TATAGCTA/iSp18/TATAGCTATA/36 TAMSp/-3'
hTelo	5'-AGGGTTAGGGTTAGGGTTAGGG-3'
BCL2	5'-AGGGGCGGGCGCGGGAGGAAGGGGGCGGGA-3'
cKIT-1	5'-AGGGAGGGCGCTGGGAGGAGGG-3'
cKIT-2	5'-CGGGCGGGCGCGAGGGAGGGG-3'
cKIT-SP	5'-GGCGAGGAGGGGCGTGGCCGGC-3'
сМҮС	5'-TGGGGAGGGTGGGGAGGGTGGGGAAGG-3'
RET	5'-AGGGGCGGGGGGGGGGGGGGGG-3'
kRAS	5'-AGGGCGGTGTGGGAAGAGGGAAGAGGGGGGAGG-3'

Table S2. Δ TM values of nucleic acid sequences upon interaction with metal complexes 1 – 12. Numerical values of Δ TM obtained by FRET melting assay of the considered nucleic acid sequences (dsDNA and G4 molecules) at 0.2 μ M in presence of 1 μ M of the tested metal complex (ratio [DNA]:[complex] = 1:5). Values are expressed as Mean ± SEM.

	dsDNA	hTelo	cKIT-1	cKIT-2	cKIT-SP	BCL2	kRAS	сМҮС	RET
1	1.75 ± 0.72	27.36 ± 0.17	28.10 ± 0.13	> 35	14.08 ± 1.14	> 35	22.30 ± 0.16	> 35	> 35
2	0.27 ± 0.36	9.32 ± 0.05	13.92 ± 0.37	16.71 ± 0.13	5.05 ± 0.98	8.18 ± 0.06	8.71 ± 0.19	12.66 ± 0.05	13.94 ± 1.39
3	1.69 ± 0.25	22.17 ± 0.10	23.46 ± 0.34	21.54 ± 0.11	11.84 ± 0.09	14.16 ± 0.02	17.61 ± 0.17	> 35	> 35
4	1.58 ± 0.06	13.40 ± 1.87	27.00 ± 0.06	18.39 ± 0.35	15.25 ± 1.60	12.00 ± 0.52	30.49 ± 0.71	19.66 ± 0.67	15.28 ± 0.18
5	0.75 ± 0.24	2.49 ± 0.03	11.39 ± 0.27	16.10 ± 0.06	10.46 ± 0.71	5.30 ± 0.26	14.58 ± 0.26	-2.11 ± 0.29	10.01 ± 0.20
6	0.15 ± 0.49	6.95 ± 0.06	8.36 ± 0.07	14.12 ± 0.03	-0.51 ± 1.07	5.61 ± 0.09	10.99 ± 0.05	13.14 ± 0.16	14.06 ± 0.19
7	0.27 ± 0.01	0.62 ± 0.13	1.09 ± 0.01	1.45 ± 0.06	1.28 ± 0.85	0.59 ± 0.04	0.79 ± 0.28	1.66 ± 0.11	1.11 ± 0.06
8	1.18 ± 0.22	2.08 ± 0.60	6.02 ± 0.01	7.11 ± 0.09	1.52 ± 0.35	2.91 ± 0.05	5.33 ± 0.18	8.74 ± 0.18	9.77 ± 0.03
9	1.22 ± 0.45	9.71 ± 0.16	11.44 ± 0.14	13.33 ± 0.11	2.09 ± 0.92	6.98 ± 0.09	12.20 ± 0.10	9.64 ± 0.03	> 35
10	1.75 ± 0.65	26.59 ± 0.16	28.94 ± 0.10	> 35	11.84 ± 0.08	> 35	24.68 ± 0.22	> 35	> 35
11	0.83 ± 0.98	9.60 ± 2.60	16.38 ± 1.37	31.00 ± 0.04	4.07 ± 2.21	7.35 ± 0.43	4.00 ± 0.10	> 35	11.05 ± 0.28
12	1.41 ± 0.11	29.91 ± 0.35	30.73 ± 0.58	> 35	9.71 ± 0.07	17.81 ± 0.10	20.21 ± 0.58	> 35	> 35

Table S3. Proliferation inhibition rate of cancer cells by metal complexes 1 - 12. Proliferation inhibition rate (%) in T3M4, T47D and MDA-MB-231 cell lines with 10 or 50 μ M of the corresponding complex for 48 or 72 hours. Proliferation inhibition of each condition was calculated relatively to its time-control and was expressed in percentage (%). *Mean* ± *SEM* (n=3).

	-	T3	M4	T4	7D	MDA-I	MB-231
	-	10 µM	50 µM	10 µM	50 µM	10 µM	50 μM
1	48 h	12.4 ± 0.8	61.9 ± 0.4	-13.6 ± 3.8	59.3 ± 1.5	-3.9 ± 5.1	32.0 ± 1.1
T	72 h	13.9 ± 3.7	83.8 ± 0.2	-0.5 ± 2.3	76.4 ± 2.2	-3.1 ± 5.9	35.4 ± 3.4
2	48 h	16.8 ± 5.6	57.6 ± 4.7	11.7 ± 2.9	62.9 ± 0.9	10.6 ± 2.4	30.9 ± 6.9
-	72 h	17.2 ± 9.1	72.9 ± 6.7	7.0 ± 2.0	86.5 ± 0.4	13.2 ± 1.5	40.3 ± 1.5
3	48 h	28.8 ± 0.5	59.0 ± 3.5	-0.1 ± 2.8	18.3 ± 1.2	-2.1 ± 1.9	5.2 ± 1.5
5	72 h	58.4 ± 1.3	87.7 ± 0.7	-2.6 ± 4.5	33.5 ± 4.6	1.1 ± 2.2	-4.5 ± 3.9
4	48 h	24.9 ± 4.8	62.0 ± 0.2	3.7 ± 1.6	81.9 ± 1.4	5.7 ± 1.1	69.0 ± 5.0
4	72 h	27.6 ± 2.7	86.7 ± 0.4	7.6 ± 1.7	97.8 ± 0.3	2.2 ± 1.9	76.2 ± 3.4
F	48 h	22.7 ± 1.6	60.6 ± 4.5	13.9 ± 3.8	62.4 ± 0.7	17.8 ± 9.7	77.3 ± 2.4
5	72 h	24.8 ± 8.6	75.7 ± 8.6	6.1 ± 3.6	87.8 ± 1.9	13.8 ± 9.1	93.9 ± 1.9
~	48 h	13.8 ± 3.3	47.6 ± 3.9	-5.9 ± 3.2	51.8 ± 3.5	7.1 ± 1.6	39.6 ± 2.9
0	72 h	14.7 ± 4.1	49.4 ± 2.4	5.0 ± 2.1	57.5 ± 3.4	7.9 ± 1.9	51.7 ± 5.7
7	48 h	15.2 ± 9.6	29.5 ± 15.0	4.9 ± 6.3	50.9 ± 3.3	18.9 ± 1.4	25.5 ± 1.4
,	72 h	17.2 ± 12.0	51.7 ± 14.0	6.3 ± 3.7	72.9 ± 1.5	19.4 ± 1.5	29.4 ± 2.6
0	48 h	6.2 ± 5.3	41.0 ± 3.2	9.9 ± 2.1	21.4 ± 3.6	11.1 ± 0.5	48.2 ± 6.1
0	72 h	13.8 ± 3.0	54.2 ± 0.4	11.0 ± 2.7	52.5 ± 4.4	21.2 ± 7.0	64.9 ± 3.7
0	48 h	13.0 ± 3.0	15.89 ± 5.84	9.4 ± 2.5	32.9 ± 0.9	17.9 ± 4.9	17.4 ± 4.9
9	72 h	18.9 ± 3.8	33.57 ± 6.48	9.3 ± 3.9	46.6 ± 2.3	16.1 ± 2.6	14.8 ± 1.2
10	48 h	20.6 ± 5.5	50.76 ± 0.28	11.1 ± 2.9	89.3 ± 0.7	2.2 ± 2.6	51.1 ± 3.3
10	72 h	19.6 ± 3.1	66.78 ± 2.36	11.9 ± 0.7	97.2 ± 0.2	-2.2 ± 3.8	54.4 ± 3.6
11	48 h	22.4 ± 12.3	43.84 ± 11.02	30.8 ± 5.0	27.4 ± 1.8	24.2 ± 4.9	38.2 ± 6.5
11	72 h	29.3 ± 15.4	65.02 ± 11.60	27.8 ± 5.5	67.5 ± 4.1	26.3 ± 1.7	54.6 ± 9.2
12	48 h	12.2 ± 1.7	45.50 ± 1.71	13.5 ± 0.2	39.7 ± 2.7	7.3 ± 2.1	26.6 ± 5.3
12	72 h	14.9 ± 1.8	58.26 ± 1.08	8.5 ± 4.2	46.6 ± 4.5	13.2 ± 7.9	28.4 ± 8.5

Table S4. RT-qPCR primers sequences (5' - 3').

GENE	Sens	Primer sequence	Temp. (°C)
IIRR	Forward	5' – GCTTTGTTGGGTGAGCTTGT – 3'	60°C
066	Reverse	5' – CGAAGATCTGCATTTTGACCT – 3'	00 C
TPD	Forward	5' – GAGCTGTGATGTGAAGTTTCC – 3'	60°C
IBP	Reverse	5' - TCTGGGTTTGATCATTCTGTAG - 3'	60 C
DDI 12A	Forward	5' - GTTCCTGCTCTCAAG – 3'	۲۹°C
RPLISA	Reverse	5' – GTCACTGCCTGGTACTTCC - 3'	56 C
LDAS	Forward	5' – TCTTGCCTCCCTACCTTCCACAT - 3'	61°C
KNAS	Reverse	5' – CTGTCAGATTCTCTTGAGCCCTG - 3'	01 C
CMAYC	Forward	5' -GGCTCCTGGCAAAAGGTCA - 3'	61°C
LIVITC	Reverse	5' – CTGCGTAGTTGTGCTGATGT - 3'	61 C
RCID	Forward	5' – GTCATGTGTGTGGAGAGCGTCAACC - 3'	60°C
BCL2	Reverse	5' – CCAGGGCCAAACTGAGCAGAGTC - 3'	00 C
PET	Forward	5' – GCAGCATTGTTGGGGGGACA - 3'	61°C
	Reverse	5' - CACCGGAAGAGGAGTAGCTG - 3'	UT C