

Supplemental Figure 4

MDA-MB-231

mRNA	Position	Complex	Heteroplasmy
Missense	A8860G	ATP8 (C-V)	
	C12084T	ND4 (C-I)	
	A13966G	ND5 (C-I)	
	C14766T	CYTB (C-III)	143/145 (98.6%)
	A15326G	CYTB (C-III)	
Frameshift			
Stop codon			
Modifiers			
D-loop	A73G		
	A153G		
	T195C		
	A263G		
	A302AC		
	T310TC		
	G513GCA		46/48 (95.8%)
	T16093C		
	C16184A		36/95 (37.9%)
	T16189C		
	C16223T		
	A16265G		
	C16278T		
T16519C			
tRNA			
rRNA	G709A		
	A750G		
	A1438G		
	G1719A		
	A2706G		
Synonymous			
	A4769G	ND2 (C-I)	
	T6221C	COX1 (C-IV)	
	C6371T	COX1 (C-IV)	
	C7028T	COX1 (C-IV)	
	T8506C	ATP6 (C-V)	
	C9099T	ATP8 (C-V)	36/224 (16.1%)
	G11719A	ND4 (C-I)	
	C12705T	ND5 (C-I)	
	T14470C	ND6 (C-I)	
	T15310C	CYTB (C-III)	