
Primer pair	Forward sequence	Reverse sequence	Target	5' end	3' end
URA3	ATGTCGAAAGCTACAT ATAAGGAACG	GGTGGTACGAACATCC AATGA	URA3	1	123
DHFR	ATGGTTCGACCATTGA ACTGC	CACTGAAGAGGTTGTG GTCATTC	mDHFR	1	133
