

Gene	Gene. ID	AChange
CYP2B6	CYP2B6:NM 000767. 4:exon4	p. Q172H (c. G516T)
CYP2B6	CYP2B6:NM 000767. 4:exon5	p. K262R (c. A785G)
CYP2D6	CYP2D6:NM 001025161. 2:exon1	p. P34S (c. C100T)
CYP3A5	.	.
ERCC1	ERCC1:NM 202001. 2:exon3	p. N118N (c. T354C)
GSTT1	.	.
MTHFR	MTHFR:NM 005957. 4:exon5	p. A222V (c. C665T)
UGT1A1	UGT1A1:NM 000463. 2	.
XRCC1	XRCC1:NM 006297. 2:exon10	p. Q399R (c. A1196G)
CDKN2A	CDKN2A:NM 000077. 4:exon2	p. G67V (c. G200T)
EGFR	EGFR:NM 005228. 3:exon19	p. 745 750del (c. 2235 2249delGGAAT)
HGF	HGF:NM_000601. 4:exon6	p. E210D (c. A630C)
TP53	TP53:NM 001126112. 2:exon7	p. R248Q (c. G743A)
EGFR	.	.
NRAS	.	.

ExonicFunc	AF
missense variant	.
missense variant	.
missense variant	.
intron variant	.
synonymous variant	.
.	.
missense variant	.
upstream gene variant	.
missense variant	.
missense variant	99.6%
inframe deletion	56.4%
missense_variant	24.1%
missense variant	99.7%
.	.
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