

**Influence of *CYP2C8\*3* and *ABCG2 C421A* genetic polymorphisms on trough concentration and molecular response of imatinib in Egyptian patients with chronic myeloid leukemia**

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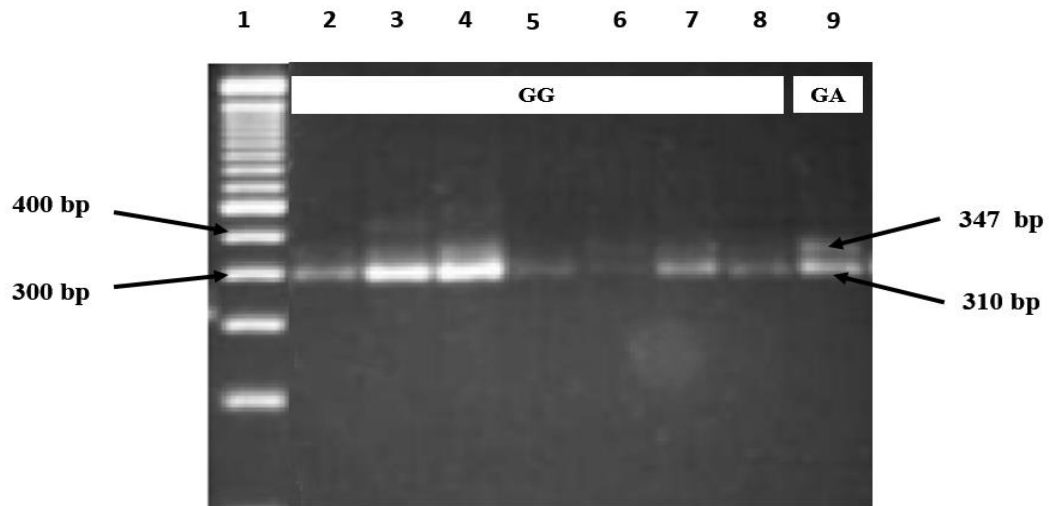
Orchid ID: <https://orcid.org/0000-0002-5990-6152>

**Details of the polymerase chain reaction-restriction fragment length polymorphism (PCR-RFLP) of *CYP2C8\*3* and *ABCG2 C421A* polymorphisms:**

- a- **Polymerase chain reaction amplification of DNA samples:** A 50  $\mu\text{L}$  reaction mixture was prepared in a 0.2 mL PCR tube following the manufacturer's protocol. The mixture included 25  $\mu\text{L}$  of COSMO PCR Master Mix, 2  $\mu\text{L}$  each of forward and reverse primers, 11  $\mu\text{L}$  of nuclease-free water, and 10  $\mu\text{L}$  of the DNA template.
- b- **Restriction fragment length polymorphism analysis:** The amplified PCR products for each gene were digested with the appropriate restriction enzymes according to the manufacturer's protocol. For each sample, a 30  $\mu\text{L}$  restriction reaction mixture was prepared, containing 17  $\mu\text{L}$  of nuclease-free water, 2  $\mu\text{L}$  of buffer, 10  $\mu\text{L}$  of PCR product, and 1  $\mu\text{L}$  of the restriction enzyme.

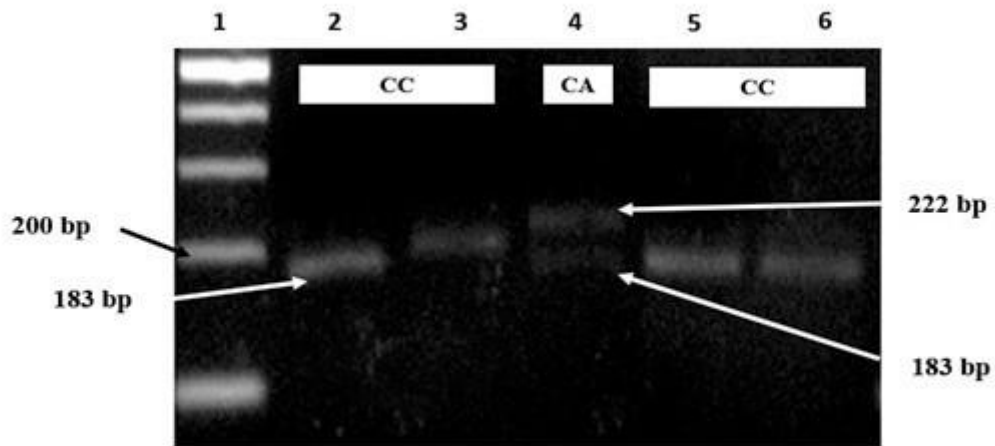
The primer sequences, PCR thermal conditions, restriction enzymes, and incubation temperatures for each SNP are presented in the following table:

	<b><i>CYP2C8*3 (G416A; rs11572080)</i></b>	<b><i>ABCG2 C421A (rs2231142)</i></b>
<b>Primer sequence:</b>		
- Forward	5' AGGCAATTCCCCAATATCTC-3'	5'- TGTTGTGATGGGCACTCTGATGT G-3'
- Reverse	5'-ACTCCTCCACAAGGCAGTGA-3'	5'- ATCAGAGTCATTTTATCCACAC - 3'
<b>PCR thermal conditions:</b>		
- Initial denaturation	94 °C for 2 minutes.	94 °C for 10 minutes.
- Denaturation	94 °C for 30 seconds	
- Annealing	48 °C for 80 seconds.	53 °C for 30 seconds
- Extension	72 °C for 1 minute.	53 °C for 30 seconds
- Number of cycles	35 cycles	
- Final extension	72 °C for 7 minutes.	
<b>PCR product [base pair (bp)]</b>	347	222
<b>Restriction enzyme</b>	<i>Bse</i> RI	<i>Hpy</i> CH4III
<b>Digestion temperature and duration</b>	37 °C for 15 minutes	37°C for 1 hours
<b>Fragments length after digestion (bp):</b>		
- Homozygous wild type	<b>GG:</b> 310.	<b>CC:</b> 183.
- Heterozygous type	<b>GA:</b> 347 and 310.	<b>CA:</b> 222 and 183



**Gel electrophoresis of *CYP2C8\*3* (G416A; rs11572080) polymorphism following PCR-RFLP analysis.**

Lane 1 contains a 100 bp DNA ladder. Lanes 2–8 display the 310 bp band, indicating the wild-type homozygous genotype (GG; *CYP2C8\*1/\*1*). Lane 9 shows the 347 bp and 310 bp bands, representing the heterozygous genotype (GA; *CYP2C8\*1/\*3*).



**Gel electrophoresis of *ABCG2* C421A (rs2231142) polymorphism following PCR-RFLP analysis.**

Lane 1 contains a 100 bp DNA ladder. Lanes 2, 3, 5, and 6 display the 183 bp band, indicating the wild-type homozygous genotype (CC). Lane 4 shows both the 222 bp and 183 bp bands, representing the heterozygous genotype (CA).