



## Specific c-Jun N-Terminal Kinase Inhibitor, JNK-IN-8 Suppresses Mesenchymal Profile of PTX-Resistant MCF-7 Cells through Modulating PI3K/Akt, MAPK and Wnt Signaling Pathways

Pelin Ozfiliz-Kilbas, Ozlem Sonmez, Pinar Uysal-Onganer, Ajda Coker-Gurkan, Pinar Obakan-Yerlikaya and Elif Damla Arisan



**Figure S1.** Densitometry analysis of PI3K/Akt (**A**) and MAPK (**B**) pathway members demonstrated in Figure 3. Statistically significant values considered as following: A) PI3K, phospho-Akt S473, phospho- GSK3 $\beta$  S9 and Cyclin D1 \*\*\*\* p < 0.0001. B)phospho-c-RAF \*\* p = 0.0049, p38 \*\* p = 0.0060, phospho-p38 \*\* p = 0.0013, p44/42 \* p= 0.0381, phospho-p44/42 \* = 0.0130, \*\* p = 0.0016, c-Jun \*\* p = 0.0023 phospho-SAPK/JNK \*\* p = 0.0013, phospho-AMPK- $\alpha$  T172 \*\*\*\* p < 0.0001.





**Figure S2.** (**A**) The mRNA expression levels of Twist in wt and PTX-res cells determined by RT-PCR. 18S was used as a loading control. (**B**) Densitometry analysis of EMT pathway members demonstrated in Figure 4. Statistically significant values considered as following: E-cadherin \*\*\*\* p < 0.0001, Claudin \*\* p = 0.0013, \*\*\*\* p < 0.0001, N-cadherin and Vimentin \*\*\*\* p < 0.0001, Snail \*\* p = 0.0084, \*\*\* p = 0.0007, TCF8/ZEB1 \*\*\* p = 0.0003,  $\beta$ -catenin \* p = 0.0088, Axin-1\*\*\*\* p < 0.0001.



**Figure S3.** Densitometry analysis of p38, phospho-p38, SAPK/JNK, phospho-SAPK/JNK, E-cadherin, Vimentin, and MDR/ABCB1 expressions demonstrated in Figure 5. Statistically significant values considered as following: p38 \*\*\*\* p < 0.0001, \*\* p = 0.0036, phospho-p38-MAPK Thr180/Tyr182 \*\* p = 0.0079, \*\*\*\* p < 0.0001, phospho-SAPK/JNK, SAPK/JNK \*\*\*\* p < 0.0001, E-cadherin \*\*\* p = 0.0001, Vimentin and MDR/ABCB1 \*\*\*\* p < 0.0001.



**Figure S4.** Densitometry analysis of Wnt signaling members demonstrated in Figure 6. Statistically significant values considered as following: Wnt3a \*\* p = 0.0010, \*\*\* p = 0.0001, Wn5a/b, phospho- LRP6 Ser1490 \*\*\*\* p < 0.0001, LRP6 \*\* p = 0.0063, \*\*\* p = 0.0004, Dvl2 \*\*p = 0.0014, \*\*\* p = 0.0002, Dvl3 \*\*\*\* p < 0.0001, Axin1 \*\*\*\* p < 0.0001, \*p = 0.0150.



© 2020 by the authors. Licensee MDPI, Basel, Switzerland. This article is an open access article distributed under the terms and conditions of the Creative Commons Attribution (CC BY) license (http://creativecommons.org/licenses/by/4.0/).