

THE STUDY OF THE RELATIONSHIP OF HS6ST2 WITH CXCL14 AND IFIT1 IN BREAST CANCER

From previous study, the silencing of *HS6ST2* in MCF7 demonstrated enhanced migration and invasion, while the DNA microarray picked out two candidate genes: *CXCL14* and *IFIT1*. Primarily, findings from *HS6ST2* silencing were corroborated via *HS6ST2* over-expression in MDA-MB-231 and MCF7, and they were in agreement. Through literature study, *CXCL14* has a role in promoting migration and invasion, indicative of *CXCL14*'s involvement in *HS6ST2*-silenced cells. Using double silencing, the invasion and migration capabilities seen in *HS6ST2*-silenced cells were abolished and the cells behaved more like non-silenced cells. Additionally, DNA microarray together with gene ontology postulated that the changes observed could be further attributed to the TGF- β pathway through SMAD2/3 signalling. On the contrary, *IFIT1* has unreported functions in breast cancer. Novel findings of *IFIT1* have shown the protein to be pro-tumour advancement. Through *IFIT1* silencing in MDA-MB-231 and MCF7, there were significant decrease in migration, invasion and proliferation. Additionally, IHC of *HS6ST2*, *CXCL14* and *IFIT1* in IDC TMAs were also performed to observe distribution in the tissues and various statistical analyses were done to evaluate the correlation of staining intensities with clinicopathological parameters, disease survival and recurrence. To date, this study is the first in exploring the possible relationship between *HS6ST2* and *CXCL14*, as well as *IFIT1*, in breast cancer progression.

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