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- 1) Ferrari, A., Vincent-Salomon, A., Pivot, X., Sertier, A.-S., Thomas, E., Tonon, L., Boyault, S., Mulugeta, E., Treilleux, I., MacGrogan, G., Arnould, L., Kielbassa, J., Le Texier, V., Blanché, H., Deleuze, J.-F., Jacquemier, J., Mathieu, M.-C., Penault-Llorca, F., Bibeau, F., Mariani, O., Mannina, C., Pierga, J.-Y., Trédan, O., Bachelot, T., Bonnefoi, H., Romieu, G., Fumoleau, P., Delaloge, S., Rios, M., Ferrero, J.-M., Tarpin, C., Bouteille, C., Calvo, F., Gut, I.G., Gut, M., Martin, S., Nik-Zainal, S., Stratton, M.R., Pauporté, I., Saintigny, P., Birnbaum, D., Viari, A., Thomas, G.
[A whole-genome sequence and transcriptome perspective on HER2-positive breast cancers](#)
(2016) Nature Communications, 7, art. no. 12222, . Cited 15 times.

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