

Table SVI. Comparative analysis with TCGA-TNBC.

Chromosomal region	Cohort (n=5)	TCGA-TNBC (n=139)	Observations
1q23.3 (gains)	100% (5/5)	23.7% (33/139)	Recurrent in the present cohort; moderate frequency in TCGA-TNBC
8q24.3 (gains, MYC)	100% (5/5)	35.3% (49/139)	Canonical TNBC alteration; consistent with published data
6q25.2 (loss, ESR1)	100% (5/5)	12.2% (17/139)	Present in all the cohort samples; less frequent in TCGA-TNBC
20q13.33 (gains)	60% (3/5)	28.1% (39/139)	Moderate frequency; comparable to TCGA-TNBC

TCGA, The Cancer Genome Atlas; TNBC, triple negative breast cancer; ESR1, estrogen receptor 1.