

S2 Table. Bonferroni correction for multiple comparisons among molecular subtypes according to biologic parameters

Subtype	RPPA				mRNA-seq				Methylation				CNA lin			
	LumA	LumB	HER2	TN	LumA	LumB	HER2	TN	LumA	LumB	HER2	TN	LumA	LumB	HER2	TN
LumA		<0.001	<0.001	<0.001		<0.001	<0.001	<0.001		<0.001	<0.001	<0.001		0.073	0.552	0.225
LumB	<0.001		<0.001	<0.001	<0.001		<0.001	<0.001	<0.001		<0.001	<0.001	0.073		1.000	1.000
HER2	<0.001	<0.001		1.000	<0.001	<0.001		1.000	<0.001	<0.001		1.000	0.552	1.000		1.000
TN	<0.001	<0.001	1.000		<0.001	<0.001	1.000		<0.001	<0.001	1.000		0.225	1.000	1.000	

RPPA, reverse phase protein array; mRNA-seq, mRNA sequencing; CNA lin, copy number alteration linear; LumA, luminal A; LumB, luminal B; HER2, human epidermal growth factor receptor 2; TN, triple negative.